

protein	annotation	FDR			
		e15.5	e17.5	p2	p14
A2ABG4	TBC1 domain family, member 16 OS=Mus musculus GN=Tbc1d16 PE=4 SV=1	1.00	1.00	1.00	1.00
A2ARZ3	Fibrous sheath-interacting protein 2 OS=Mus musculus GN=Fspip2 PE=1 SV=3	1.00	0.05	1.00	1.00
A2ASQ1	Agriin OS=Mus musculus GN=Agriin PE=2 SV=1	0.07	1.00	1.00	1.00
A8DUK4	Beta-globin OS=Mus musculus GN=Hbb-b1 PE=3 SV=1	1.00	0.00	0.00	0.00
B2RS76	Carboxypeptidase B1 (Tissue) OS=Mus musculus GN=Cpb1 PE=2 SV=1	1.00	0.00	0.00	0.00
B2RX14	Terminal uridylyltransferase 4 OS=Mus musculus GN=Zcchc11 PE=1 SV=2	1.00	1.00	1.00	0.06
B7ZNL3	Tpm1 protein OS=Mus musculus GN=Tpm1 PE=2 SV=1	1.00	0.03	0.02	0.02
D3YY62	Uncharacterized protein OS=Mus musculus GN=Gm9858 PE=4 SV=1	0.03	0.03	0.02	1.00
D3Z5Y1	Uncharacterized protein OS=Mus musculus GN=Stard13 PE=4 SV=1	0.03	1.00	1.00	1.00
D3Z6P0	Protein disulfide-isomerase A2 OS=Mus musculus GN=Pdia2 PE=1 SV=1	1.00	0.02	0.00	0.00
E9PUV7	Uncharacterized protein OS=Mus musculus GN=4930485B16Rik PE=4 SV=1	1.00	0.08	1.00	1.00
E9PVM7	Uncharacterized protein OS=Mus musculus GN=Gstm5 PE=3 SV=1	1.00	1.00	1.00	0.03
E9PVZ3	Uncharacterized protein OS=Mus musculus GN=Ccdc144b PE=4 SV=1	1.00	1.00	1.00	0.04
E9PZQ0	Uncharacterized protein OS=Mus musculus GN=Ryr1 PE=4 SV=1	1.00	0.07	1.00	0.04
E9Q191	Uncharacterized protein OS=Mus musculus GN=Oifr288 PE=3 SV=1	1.00	1.00	0.08	1.00
E9Q1Y9	Uncharacterized protein OS=Mus musculus GN=5430421N21Rik PE=3 SV=1	1.00	0.00	1.00	1.00
E9Q447	Uncharacterized protein OS=Mus musculus GN=Spna2 PE=4 SV=1	1.00	0.00	0.00	0.00
E9Q455	Uncharacterized protein OS=Mus musculus GN=Tpm1 PE=3 SV=1	1.00	0.03	0.02	0.00
E9Q557	Desmoplakin OS=Mus musculus GN=Dsp PE=3 SV=1	1.00	0.00	0.00	0.00
E9Q616	Uncharacterized protein OS=Mus musculus GN=Ahnak PE=4 SV=1	1.00	0.00	0.00	0.00
E9Q7L1	Uncharacterized protein OS=Mus musculus GN=Urb2 PE=4 SV=1	0.01	0.01	0.02	1.00
E9Q9C6	Uncharacterized protein OS=Mus musculus GN=Fgcbp PE=4 SV=1	1.00	0.03	1.00	1.00
F6Q3G7	Uncharacterized protein (Fragment) OS=Mus musculus GN=Upf2 PE=4 SV=1	0.03	0.03	1.00	1.00
F6R5A0	Uncharacterized protein (Fragment) OS=Mus musculus GN=D1Ert622e PE=4 SV=1	1.00	0.04	1.00	1.00
F6VBB8	Uncharacterized protein (Fragment) OS=Mus musculus PE=4 SV=1	0.02	0.03	0.03	1.00
F6W6S4	Uncharacterized protein (Fragment) OS=Mus musculus GN=Hpn PE=4 SV=1	1.00	1.00	0.04	1.00
F7DBB3	Uncharacterized protein (Fragment) OS=Mus musculus GN=Ahnak2 PE=4 SV=1	1.00	1.00	1.00	0.00
O08547	Vesicle-trafficking protein SEC22b OS=Mus musculus GN=Sec22b PE=1 SV=3	1.00	1.00	1.00	0.02
O08553	Dihydropyrimidinase-related protein 2 OS=Mus musculus GN=Dpysl2 PE=1 SV=2	1.00	0.02	0.02	0.00
O08749	Dihydrolipoyl dehydrogenase, mitochondrial OS=Mus musculus GN=Dld PE=1 SV=2	1.00	1.00	0.08	1.00
O08795	Glucosidase 2 subunit beta OS=Mus musculus GN=Prksh PE=1 SV=1	0.02	0.00	1.00	0.06
O08807	Peroxisomal protein OS=Mus musculus GN=Prdx4 PE=1 SV=1	1.00	0.00	0.02	1.00
O35074	Prostacyclin synthase OS=Mus musculus GN=Ptgis PE=2 SV=1	1.00	1.00	1.00	0.00
O35098	Dihydropyrimidinase-related protein 4 OS=Mus musculus GN=Dpysl4 PE=1 SV=1	1.00	1.00	1.00	0.02
O35129	Prohibitin-2 OS=Mus musculus GN=Phb2 PE=1 SV=1	1.00	0.02	0.02	1.00
O35326	Serine/arginine-rich splicing factor 5 OS=Mus musculus GN=Srsf5 PE=1 SV=1	1.00	1.00	0.02	1.00
O35350	Calpain-1 catalytic subunit OS=Mus musculus GN=Capn1 PE=2 SV=1	1.00	1.00	1.00	0.07
O35381	Acidic leucine-rich nuclear phosphoprotein 32 family member A OS=Mus musculus GN=Anp32a PE=1 SV=1	1.00	0.00	1.00	1.00
O35639	Annexin A3 OS=Mus musculus GN=Anxa3 PE=1 SV=4	1.00	1.00	1.00	0.00
O35640	Annexin A8 OS=Mus musculus GN=Anxa8 PE=2 SV=2	1.00	1.00	1.00	0.05
O35682	Myeloid-associated differentiation marker OS=Mus musculus GN=Myadm PE=2 SV=2	1.00	1.00	1.00	0.02
O35685	Nuclear migration protein nudC OS=Mus musculus GN=Nudc PE=1 SV=1	1.00	0.04	1.00	1.00
O35855	Branched-chain-amino-acid aminotransferase, mitochondrial OS=Mus musculus GN=Bcat2 PE=2 SV=2	1.00	0.03	1.00	1.00
O35887	Calumenin OS=Mus musculus GN=Calu PE=1 SV=1	1.00	0.00	0.00	0.00
O54724	Polymerase I and transcript release factor OS=Mus musculus GN=Ptrf PE=1 SV=1	1.00	1.00	0.02	0.00
O54734	Dolichyl-diphosphooligosaccharide--protein glycosyltransferase 48 kDa subunit OS=Mus musculus GN=Ddost PE=1 SV=1	1.00	0.02	1.00	0.00
O54754	Aldehyde oxidase OS=Mus musculus GN=Aox1 PE=2 SV=2	0.04	1.00	1.00	1.00
O54879	High mobility group protein B3 OS=Mus musculus GN=Hmgb3 PE=2 SV=3	1.00	1.00	0.02	1.00
O54962	Barrier-to-autointegration factor OS=Mus musculus GN=Banf1 PE=1 SV=1	0.02	1.00	1.00	1.00
O55022	Membrane-associated progesterone receptor component 1 OS=Mus musculus GN=Pgrmc1 PE=1 SV=4	1.00	0.02	1.00	1.00
O55023	Inositol monophosphatase 1 OS=Mus musculus GN=Impa1 PE=1 SV=1	1.00	1.00	1.00	0.07
O55026	Ectonucleoside triphosphate diphosphohydrolase 2 OS=Mus musculus GN=Entpd2 PE=1 SV=2	1.00	1.00	1.00	0.00
O55029	Coatomer subunit beta' OS=Mus musculus GN=Copb2 PE=2 SV=2	1.00	1.00	1.00	0.02
O55131	Septin-7 OS=Mus musculus GN=Sept7 PE=1 SV=1	1.00	1.00	0.00	0.02
O55142	60S ribosomal protein L35a OS=Mus musculus GN=Rpl35a PE=2 SV=2	1.00	1.00	0.02	0.02
O55234	Proteasome subunit beta type-5 OS=Mus musculus GN=Psb5 PE=1 SV=3	1.00	0.02	0.00	1.00
O70251	Elongation factor 1-beta OS=Mus musculus GN=Eef1b PE=1 SV=5	0.02	0.00	1.00	0.00
O70318	Band 4.1-like protein 2 OS=Mus musculus GN=Epb41l2 PE=1 SV=1	1.00	0.05	1.00	1.00
O70435	Proteasome subunit alpha type-3 OS=Mus musculus GN=Psa3 PE=1 SV=3	1.00	0.02	0.02	1.00
O70456	14-3-3 protein sigma OS=Mus musculus GN=Sfn PE=1 SV=2	1.00	1.00	0.02	1.00
O70475	UDP-glucose 6-dehydrogenase OS=Mus musculus GN=Ugdh PE=1 SV=1	1.00	0.00	0.00	0.00
O70492	Sorting nexin-3 OS=Mus musculus GN=Snx3 PE=1 SV=3	1.00	1.00	0.05	1.00
O88342	WD repeat-containing protein 1 OS=Mus musculus GN=Wdr1 PE=1 SV=3	1.00	1.00	0.06	1.00
O88456	Calpain small subunit 1 OS=Mus musculus GN=Capns1 PE=2 SV=1	1.00	1.00	1.00	0.02
O88487	Cytoplasmic dynein 1 intermediate chain 2 OS=Mus musculus GN=Dync1i2 PE=2 SV=1	1.00	0.00	1.00	1.00
O88569	Heterogeneous nuclear ribonucleoproteins A2/B1 OS=Mus musculus GN=Hnmpa2b1 PE=1 SV=2	1.00	0.00	0.00	0.00
O88844	Isocitrate dehydrogenase [NADP] cytoplasmic OS=Mus musculus GN=Idh1 PE=1 SV=2	1.00	0.00	0.00	0.00
O88962	7-alpha-hydroxycholesterol-4-en-3-one 12-alpha-hydroxylase OS=Mus musculus GN=Cyp8b1 PE=2 SV=1	1.00	1.00	1.00	0.07
O89086	Putative RNA-binding protein 3 OS=Mus musculus GN=Rbm3 PE=1 SV=1	0.03	0.02	0.02	1.00

P00683	Ribonuclease pancreatic OS=Mus musculus GN=Rnase1 PE=1 SV=2	1.00	0.02	0.02	1.00
P00688	Pancreatic alpha-amylase OS=Mus musculus GN=Amy2 PE=1 SV=2	1.00	0.00	0.00	0.00
P04104	Keratin, type II cytoskeletal 1 OS=Mus musculus GN=Krt1 PE=1 SV=4	1.00	0.02	1.00	0.02
P04117	Fatty acid-binding protein, adipocyte OS=Mus musculus GN=Fabp4 PE=1 SV=3	1.00	1.00	1.00	0.00
P05064	Fructose-bisphosphate aldolase A OS=Mus musculus GN=Aldoa PE=1 SV=2	0.02	0.00	0.00	0.00
P05202	Aspartate aminotransferase, mitochondrial OS=Mus musculus GN=Got2 PE=1 SV=1	1.00	1.00	0.02	1.00
P05208	Chymotrypsin-like elastase family member 2A OS=Mus musculus GN=Cela2a PE=2 SV=1	1.00	0.00	0.00	0.00
P05213	Tubulin alpha-1B chain OS=Mus musculus GN=Tuba1b PE=1 SV=2	0.00	1.00	1.00	0.00
P06151	L-lactate dehydrogenase A chain OS=Mus musculus GN=Ldha PE=1 SV=3	0.02	0.00	0.00	0.00
P06745	Glucose-6-phosphate isomerase OS=Mus musculus GN=Gpi PE=1 SV=4	1.00	1.00	0.00	0.02
P07146	Anionic trypsin-2 OS=Mus musculus GN=Prss2 PE=2 SV=1	1.00	0.00	0.02	0.00
P07356	Annexin A2 OS=Mus musculus GN=Anxa2 PE=1 SV=2	0.00	0.00	0.00	0.00
P07724	Serum albumin OS=Mus musculus GN=Alb PE=1 SV=3	1.00	0.02	1.00	0.00
P07901	Heat shock protein HSP 90-alpha OS=Mus musculus GN=Hsp90aa1 PE=1 SV=4	0.00	0.00	0.00	0.00
P08003	Protein disulfide-isomerase A4 OS=Mus musculus GN=Pdia4 PE=1 SV=3	0.00	0.00	0.00	0.00
P08113	Endoplasmin OS=Mus musculus GN=Hsp90b1 PE=1 SV=2	0.00	0.00	0.00	0.00
P08207	Protein S100-A10 OS=Mus musculus GN=S100a10 PE=2 SV=2	1.00	0.02	0.02	0.00
P08228	Superoxide dismutase [Cu-Zn] OS=Mus musculus GN=Sod1 PE=1 SV=2	0.02	0.00	0.00	0.00
P08249	Malate dehydrogenase, mitochondrial OS=Mus musculus GN=Mdh2 PE=1 SV=3	1.00	0.00	0.00	0.00
P08730	Keratin, type I cytoskeletal 13 OS=Mus musculus GN=Krt13 PE=1 SV=2	0.03	1.00	1.00	0.02
P08752	Guanine nucleotide-binding protein (G(i) subunit alpha-2 OS=Mus musculus GN=Gnai2 PE=1 SV=5	1.00	1.00	1.00	0.02
P09055	Integrin beta-1 OS=Mus musculus GN=Itgb1 PE=1 SV=1	1.00	0.03	0.08	0.04
P09103	Protein disulfide-isomerase OS=Mus musculus GN=P4hb PE=1 SV=2	0.00	0.00	0.00	0.00
P09405	Nucleolin OS=Mus musculus GN=Ncl PE=1 SV=2	0.00	0.00	0.00	0.00
P09411	Phosphoglycerate kinase 1 OS=Mus musculus GN=Pgk1 PE=1 SV=4	1.00	0.08	1.00	0.03
P09602	Non-histone chromosomal protein HMG-17 OS=Mus musculus GN=Hmgn2 PE=1 SV=2	1.00	0.00	1.00	1.00
P10107	Annexin A1 OS=Mus musculus GN=Anxa1 PE=1 SV=2	1.00	1.00	0.00	0.00
P10126	Elongation factor 1-alpha 1 OS=Mus musculus GN=Eef1a1 PE=1 SV=3	0.00	0.00	0.00	0.00
P10639	Thioredoxin OS=Mus musculus GN=Txn PE=1 SV=3	0.02	0.00	0.00	0.00
P10649	Glutathione S-transferase Mu 1 OS=Mus musculus GN=Gstm1 PE=1 SV=2	1.00	0.00	0.00	0.00
P10922	Histone H1.0 OS=Mus musculus GN=H1f0 PE=2 SV=4	0.02	0.00	0.00	0.00
P11031	Activated RNA polymerase II transcriptional coactivator p15 OS=Mus musculus GN=Sub1 PE=1 SV=3	0.02	0.00	0.02	1.00
P11499	Heat shock protein HSP 90-beta OS=Mus musculus GN=Hsp90ab1 PE=1 SV=3	0.05	0.00	0.00	0.00
P11688	Integrin alpha-5 OS=Mus musculus GN=Itga5 PE=1 SV=3	1.00	0.04	0.02	1.00
P12970	60S ribosomal protein L7a OS=Mus musculus GN=Rpl7a PE=2 SV=2	1.00	0.00	0.02	0.00
P13020	Gelsolin OS=Mus musculus GN=Gsn PE=1 SV=3	0.05	0.00	0.00	0.00
P14069	Protein S100-A6 OS=Mus musculus GN=S100a6 PE=1 SV=3	0.04	0.02	0.02	0.03
P14115	60S ribosomal protein L27a OS=Mus musculus GN=Rpl27a PE=2 SV=5	1.00	0.02	1.00	0.02
P14131	40S ribosomal protein S16 OS=Mus musculus GN=Rps16 PE=2 SV=4	1.00	0.00	0.02	0.05
P14148	60S ribosomal protein L7 OS=Mus musculus GN=Rpl7 PE=2 SV=2	1.00	0.06	1.00	0.02
P14152	Malate dehydrogenase, cytoplasmic OS=Mus musculus GN=Mdh1 PE=1 SV=3	0.02	0.00	0.00	0.00
P14206	40S ribosomal protein SA OS=Mus musculus GN=Rpsa PE=1 SV=4	0.00	0.00	0.00	0.00
P14211	Calreticulin OS=Mus musculus GN=Calr PE=1 SV=1	0.00	0.00	0.00	0.00
P14431	H-2 class I histocompatibility antigen, Q9 alpha chain (Fragment) OS=Mus musculus GN=H2-Q9 PE=2 SV=1	1.00	0.04	1.00	1.00
P14733	Lamin-B1 OS=Mus musculus GN=Lmnb1 PE=1 SV=3	0.00	0.00	0.00	0.00
P14824	Annexin A6 OS=Mus musculus GN=Anxa6 PE=1 SV=3	0.03	0.00	0.00	0.00
P14869	60S acidic ribosomal protein P0 OS=Mus musculus GN=Rplp0 PE=1 SV=3	0.02	0.00	0.02	0.00
P15626	Glutathione S-transferase Mu 2 OS=Mus musculus GN=Gstm2 PE=1 SV=2	1.00	0.02	0.00	0.00
P15864	Histone H1.2 OS=Mus musculus GN=Hist1h1c PE=1 SV=2	0.00	0.00	0.00	0.00
P15947	Kallikrein-1 OS=Mus musculus GN=Klk1 PE=1 SV=3	1.00	1.00	0.02	1.00
P16045	Galectin-1 OS=Mus musculus GN=Lgals1 PE=1 SV=3	0.02	0.00	0.00	0.00
P16858	Glyceraldehyde-3-phosphate dehydrogenase OS=Mus musculus GN=Gapdh PE=1 SV=2	0.02	0.00	0.02	0.00
P17047	Lysosome-associated membrane glycoprotein 2 OS=Mus musculus GN=Lamp2 PE=2 SV=2	1.00	1.00	0.07	1.00
P17182	Alpha-enolase OS=Mus musculus GN=Eno1 PE=1 SV=3	0.00	0.00	0.00	0.00
P17742	Peptidyl-prolyl cis-trans isomerase A OS=Mus musculus GN=Ppia PE=1 SV=2	0.06	0.00	0.00	0.00
P17751	Triosephosphate isomerase OS=Mus musculus GN=Tpi1 PE=1 SV=3	0.00	0.00	0.00	0.00
P17897	Lysozyme C-1 OS=Mus musculus GN=Lyz1 PE=1 SV=1	1.00	1.00	1.00	0.03
P17918	Proliferating cell nuclear antigen OS=Mus musculus GN=Pcna PE=1 SV=2	0.02	0.00	0.02	1.00
P18242	Cathepsin D OS=Mus musculus GN=Ctsd PE=1 SV=1	1.00	1.00	1.00	0.02
P18572	Basigin OS=Mus musculus GN=Bsg PE=1 SV=2	1.00	1.00	0.03	0.04
P18760	Cofilin-1 OS=Mus musculus GN=Cfl1 PE=1 SV=3	0.00	0.00	0.02	0.00
P19001	Keratin, type I cytoskeletal 19 OS=Mus musculus GN=Krt19 PE=2 SV=1	1.00	0.06	1.00	1.00
P19157	Glutathione S-transferase P 1 OS=Mus musculus GN=Gstp1 PE=1 SV=2	1.00	0.02	1.00	0.00
P19253	60S ribosomal protein L13a OS=Mus musculus GN=Rpl13a PE=1 SV=4	1.00	0.00	0.02	0.00
P19324	Serpin H1 OS=Mus musculus GN=Serpinh1 PE=1 SV=3	0.00	0.00	0.00	0.00
P20029	78 kDa glucose-regulated protein OS=Mus musculus GN=Hspa5 PE=1 SV=3	0.00	0.00	0.00	0.00
P20065	Thymosin beta-4 OS=Mus musculus GN=Tmsb4x PE=1 SV=1	1.00	1.00	0.06	1.00
P20108	Thioredoxin-dependent peroxide reductase, mitochondrial OS=Mus musculus GN=Prdx3 PE=1 SV=1	1.00	1.00	0.04	1.00
P20152	Vimentin OS=Mus musculus GN=Vim PE=1 SV=3	0.02	0.00	0.00	0.00
P21981	Protein-glutamine gamma-glutamyltransferase 2 OS=Mus musculus GN=Tgm2 PE=1 SV=4	1.00	1.00	1.00	0.00
P23198	Chromobox protein homolog 3 OS=Mus musculus GN=Cbx3 PE=1 SV=2	1.00	0.00	0.00	0.02

P23492	Purine nucleoside phosphorylase OS=Mus musculus GN=Pnp PE=1 SV=2	1.00	0.07	0.03	0.03
P24270	Catalase OS=Mus musculus GN=Cat PE=1 SV=4	1.00	0.02	1.00	1.00
P24369	Peptidyl-prolyl cis-trans isomerase B OS=Mus musculus GN=Ppip PE=2 SV=2	0.03	0.00	0.00	0.00
P25206	DNA replication licensing factor MCM3 OS=Mus musculus GN=Mcm3 PE=1 SV=2	0.02	1.00	1.00	1.00
P25444	40S ribosomal protein S2 OS=Mus musculus GN=Rps2 PE=1 SV=3	1.00	0.02	0.00	0.00
P26039	Talin-1 OS=Mus musculus GN=Tln1 PE=1 SV=2	1.00	0.02	0.02	0.00
P26040	Ezrin OS=Mus musculus GN=Ezr PE=1 SV=3	1.00	0.02	0.02	0.00
P26041	Moesin OS=Mus musculus GN=Msn PE=1 SV=3	0.03	0.00	0.00	0.00
P26043	Radixin OS=Mus musculus GN=Rdx PE=1 SV=3	1.00	0.00	1.00	1.00
P26350	Prothymosin alpha OS=Mus musculus GN=Ptma PE=1 SV=2	0.00	0.00	0.00	0.00
P26443	Glutamate dehydrogenase 1, mitochondrial OS=Mus musculus GN=Glud1 PE=1 SV=1	1.00	1.00	0.02	1.00
P26638	Seryl-tRNA synthetase, cytoplasmic OS=Mus musculus GN=Sars PE=2 SV=3	1.00	0.06	1.00	1.00
P26645	Myristoylated alanine-rich C-kinase substrate OS=Mus musculus GN=Marcks PE=1 SV=2	0.00	0.00	0.00	0.00
P27546	Microtubule-associated protein 4 OS=Mus musculus GN=Map4 PE=1 SV=3	1.00	0.02	0.00	0.00
P27659	60S ribosomal protein L3 OS=Mus musculus GN=Rpl3 PE=2 SV=3	1.00	0.03	0.02	0.00
P27773	Protein disulfide-isomerase A3 OS=Mus musculus GN=Pdia3 PE=1 SV=2	0.00	0.00	0.00	0.00
P28352	DNA-(apurinic or apyrimidinic site) lyase OS=Mus musculus GN=Apex1 PE=1 SV=2	1.00	1.00	0.02	1.00
P28656	Nucleosome assembly protein 1-like 1 OS=Mus musculus GN=Nap111 PE=1 SV=2	0.04	1.00	1.00	1.00
P28660	Nck-associated protein 1 OS=Mus musculus GN=Nckap1 PE=2 SV=2	1.00	0.01	1.00	1.00
P28667	MARCKS-related protein OS=Mus musculus GN=Marcks1 PE=1 SV=2	0.00	0.00	0.02	1.00
P28843	Dipeptidyl peptidase 4 OS=Mus musculus GN=Dpp4 PE=1 SV=3	1.00	1.00	1.00	0.02
P29341	Polyadenylate-binding protein 1 OS=Mus musculus GN=Pabpc1 PE=1 SV=2	0.04	1.00	0.03	1.00
P29758	Ornithine aminotransferase, mitochondrial OS=Mus musculus GN=Oat PE=1 SV=1	1.00	0.04	0.00	0.00
P30412	Peptidyl-prolyl cis-trans isomerase C OS=Mus musculus GN=Ppic PE=1 SV=1	1.00	0.05	1.00	0.02
P30416	Peptidyl-prolyl cis-trans isomerase FKBP4 OS=Mus musculus GN=Fkbp4 PE=1 SV=5	1.00	0.02	0.02	1.00
P30681	High mobility group protein B2 OS=Mus musculus GN=Hmgb2 PE=1 SV=3	0.04	0.00	0.00	0.07
P31001	Desmin OS=Mus musculus GN=Des PE=1 SV=3	1.00	0.02	0.00	0.00
P31786	Acyl-CoA-binding protein OS=Mus musculus GN=Dbi PE=1 SV=2	1.00	1.00	0.04	1.00
P34022	Ran-specific GTPase-activating protein OS=Mus musculus GN=Ranbp1 PE=1 SV=2	0.00	0.02	0.04	0.03
P34884	Macrophage migration inhibitory factor OS=Mus musculus GN=Mif PE=1 SV=2	0.04	0.02	1.00	1.00
P35278	Ras-related protein Rab-5C OS=Mus musculus GN=Rab5c PE=1 SV=2	1.00	1.00	1.00	0.00
P35486	Pyruvate dehydrogenase E1 component subunit alpha, somatic form, mitochondrial OS=Mus musculus GN=PdhE1	1.00	0.02	1.00	1.00
P35492	Histidine ammonia-lyase OS=Mus musculus GN=Hal PE=1 SV=1	1.00	1.00	1.00	0.04
P35564	Calnexin OS=Mus musculus GN=Canx PE=1 SV=1	1.00	0.00	1.00	0.00
P35700	Peroxiredoxin-1 OS=Mus musculus GN=Prdx1 PE=1 SV=1	0.00	0.00	0.00	0.00
P35979	60S ribosomal protein L12 OS=Mus musculus GN=Rpl12 PE=1 SV=2	1.00	0.00	0.00	0.00
P35980	60S ribosomal protein L18 OS=Mus musculus GN=Rpl18 PE=2 SV=3	1.00	0.00	0.00	0.00
P37804	Transgelin OS=Mus musculus GN=Tagln PE=1 SV=3	1.00	0.02	0.00	0.00
P38060	Hydroxymethylglutaryl-CoA lyase, mitochondrial OS=Mus musculus GN=Hmgcl PE=1 SV=2	1.00	1.00	0.08	1.00
P40124	Adenylyl cyclase-associated protein 1 OS=Mus musculus GN=Cap1 PE=1 SV=4	1.00	0.06	0.02	0.02
P40142	Transketolase OS=Mus musculus GN=Tkt PE=1 SV=1	0.02	0.00	0.00	0.00
P42208	Septin-2 OS=Mus musculus GN=Sept2 PE=1 SV=2	1.00	0.05	1.00	0.00
P42932	T-complex protein 1 subunit theta OS=Mus musculus GN=Cct8 PE=1 SV=3	1.00	0.02	1.00	0.05
P43137	Lithostathine-1 OS=Mus musculus GN=Reg1 PE=2 SV=1	1.00	1.00	0.00	0.00
P43274	Histone H1.4 OS=Mus musculus GN=Hist1h1e PE=1 SV=2	1.00	0.00	0.02	0.00
P43275	Histone H1.1 OS=Mus musculus GN=Hist1h1a PE=2 SV=2	0.00	0.00	0.00	0.00
P43276	Histone H1.5 OS=Mus musculus GN=Hist1h1b PE=1 SV=2	0.00	0.00	0.00	0.00
P43277	Histone H1.3 OS=Mus musculus GN=Hist1h1d PE=1 SV=2	1.00	0.05	1.00	1.00
P45376	Aldose reductase OS=Mus musculus GN=Akr1b1 PE=1 SV=3	1.00	1.00	1.00	0.02
P45591	Cofilin-2 OS=Mus musculus GN=Cfl2 PE=1 SV=1	1.00	1.00	1.00	0.02
P45878	Peptidyl-prolyl cis-trans isomerase FKBP2 OS=Mus musculus GN=Fkbp2 PE=1 SV=1	1.00	0.02	1.00	1.00
P47738	Aldehyde dehydrogenase, mitochondrial OS=Mus musculus GN=Aldh2 PE=1 SV=1	1.00	0.00	0.00	0.00
P47740	Fatty aldehyde dehydrogenase OS=Mus musculus GN=Aldh3a2 PE=2 SV=2	1.00	1.00	1.00	0.02
P47753	F-actin-capping protein subunit alpha-1 OS=Mus musculus GN=Capza1 PE=1 SV=4	1.00	0.02	0.02	0.02
P47757	F-actin-capping protein subunit beta OS=Mus musculus GN=Capzb PE=1 SV=3	1.00	0.03	0.03	0.02
P47878	Insulin-like growth factor-binding protein 3 OS=Mus musculus GN=Igfbp3 PE=2 SV=1	1.00	0.03	1.00	1.00
P47911	60S ribosomal protein L6 OS=Mus musculus GN=Rpl6 PE=1 SV=3	1.00	1.00	0.02	0.00
P47915	60S ribosomal protein L29 OS=Mus musculus GN=Rpl29 PE=2 SV=2	1.00	0.02	1.00	1.00
P47955	60S acidic ribosomal protein P1 OS=Mus musculus GN=Rplp1 PE=1 SV=1	1.00	0.02	1.00	0.02
P47962	60S ribosomal protein L5 OS=Mus musculus GN=Rpl5 PE=1 SV=3	0.03	0.00	0.00	0.08
P47963	60S ribosomal protein L13 OS=Mus musculus GN=Rpl13 PE=2 SV=3	1.00	0.00	0.00	0.04
P47964	60S ribosomal protein L36 OS=Mus musculus GN=Rpl36 PE=2 SV=2	1.00	0.00	0.04	1.00
P48036	Annexin A5 OS=Mus musculus GN=Anxa5 PE=1 SV=1	0.02	0.00	0.00	0.00
P48678	Prelamin-A/C OS=Mus musculus GN=Lmna PE=1 SV=2	0.00	0.00	0.00	0.00
P48754	Breast cancer type 1 susceptibility protein homolog OS=Mus musculus GN=Brca1 PE=1 SV=2	1.00	0.04	1.00	1.00
P48758	Carbonyl reductase [NADPH] 1 OS=Mus musculus GN=Cbr1 PE=1 SV=3	1.00	0.02	1.00	1.00
P48771	Cytochrome c oxidase subunit 7A2, mitochondrial OS=Mus musculus GN=Cox7a2 PE=1 SV=2	1.00	0.02	1.00	1.00
P49183	Deoxyribonuclease-1 OS=Mus musculus GN=Dnase1 PE=2 SV=2	0.02	0.02	1.00	1.00
P49312	Heterogeneous nuclear ribonucleoprotein A1 OS=Mus musculus GN=Hnmpa1 PE=1 SV=2	0.00	0.00	0.00	0.00
P49722	Proteasome subunit alpha type-2 OS=Mus musculus GN=Pasma2 PE=1 SV=2	1.00	0.08	1.00	1.00
P49817	Caveolin-1 OS=Mus musculus GN=Cav1 PE=1 SV=1	1.00	1.00	0.05	0.00

P49919	Cyclin-dependent kinase inhibitor 1C OS=Mus musculus GN=Cdkn1c PE=2 SV=1	1.00	0.02	1.00	1.00
P50247	Adenosylhomocysteinase OS=Mus musculus GN=Ahcy PE=1 SV=3	0.05	0.00	0.00	0.00
P50543	Protein S100-A11 OS=Mus musculus GN=S100a11 PE=2 SV=1	1.00	0.02	1.00	0.02
P50580	Proliferation-associated protein 2G4 OS=Mus musculus GN=Pa2g4 PE=1 SV=3	0.00	0.00	0.00	1.00
P51150	Ras-related protein Rab-7a OS=Mus musculus GN=Rab7a PE=1 SV=2	1.00	0.07	0.02	0.00
P51174	Long-chain specific acyl-CoA dehydrogenase, mitochondrial OS=Mus musculus GN=Acadl PE=2 SV=2	1.00	1.00	0.02	0.02
P51410	60S ribosomal protein L9 OS=Mus musculus GN=Rpl9 PE=2 SV=2	1.00	1.00	1.00	0.00
P51660	Peroxisomal multifunctional enzyme type 2 OS=Mus musculus GN=Hsd17b4 PE=1 SV=3	1.00	1.00	0.03	1.00
P51859	Hepatoma-derived growth factor OS=Mus musculus GN=Hdgf PE=1 SV=2	1.00	0.00	0.02	0.04
P52480	Pyruvate kinase isozymes M1/M2 OS=Mus musculus GN=Pkm2 PE=1 SV=4	0.00	0.00	0.00	0.00
P52793	Ephrin-A1 OS=Mus musculus GN=Efna1 PE=2 SV=1	1.00	0.07	1.00	1.00
P52927	High mobility group protein HMGI-C OS=Mus musculus GN=Hmga2 PE=1 SV=1	0.02	0.05	1.00	1.00
P53026	60S ribosomal protein L10a OS=Mus musculus GN=Rpl10a PE=1 SV=3	1.00	0.03	0.02	0.05
P54071	Isocitrate dehydrogenase [NADP], mitochondrial OS=Mus musculus GN=Idh2 PE=1 SV=3	1.00	0.00	0.00	0.00
P54227	Stathmin OS=Mus musculus GN=Stmn1 PE=1 SV=2	0.05	0.00	1.00	1.00
P56395	Cytochrome b5 OS=Mus musculus GN=Cyb5a PE=1 SV=2	1.00	1.00	1.00	0.02
P56480	ATP synthase subunit beta, mitochondrial OS=Mus musculus GN=Atp5b PE=1 SV=2	0.00	0.00	0.00	0.00
P56501	Mitochondrial uncoupling protein 3 OS=Mus musculus GN=Ucp3 PE=2 SV=1	0.05	0.03	1.00	0.08
P56959	RNA-binding protein FUS OS=Mus musculus GN=Fus PE=2 SV=1	0.02	0.00	0.00	0.00
P57759	Endoplasmic reticulum resident protein 29 OS=Mus musculus GN=Erp29 PE=1 SV=2	1.00	0.05	1.00	1.00
P57776	Elongation factor 1-delta OS=Mus musculus GN=Eef1d PE=1 SV=3	1.00	0.02	0.02	0.00
P57780	Alpha-actinin-4 OS=Mus musculus GN=Actn4 PE=1 SV=1	1.00	0.05	0.02	0.00
P58252	Elongation factor 2 OS=Mus musculus GN=Eef2 PE=1 SV=2	0.02	0.00	0.00	0.00
P58771	Tropomyosin alpha-1 chain OS=Mus musculus GN=Tpm1 PE=1 SV=1	0.02	0.00	0.00	0.00
P58871	182 kDa tankyrase-1-binding protein OS=Mus musculus GN=Tnks1bp1 PE=1 SV=2	1.00	1.00	1.00	0.06
P59158	Solute carrier family 12 member 3 OS=Mus musculus GN=Slc12a3 PE=2 SV=1	1.00	0.04	1.00	1.00
P60122	RuvB-like 1 OS=Mus musculus GN=Ruvbl1 PE=1 SV=1	1.00	1.00	0.02	1.00
P60335	Poly(rC)-binding protein 1 OS=Mus musculus GN=Pcbp1 PE=1 SV=1	1.00	0.02	0.02	0.02
P60766	Cell division control protein 42 homolog OS=Mus musculus GN=Cdc42 PE=1 SV=2	1.00	0.00	0.02	1.00
P60824	Cold-inducible RNA-binding protein OS=Mus musculus GN=Cirbp PE=1 SV=1	1.00	1.00	1.00	0.05
P60843	Eukaryotic initiation factor 4A-1 OS=Mus musculus GN=Eif4a1 PE=2 SV=1	0.03	0.00	0.02	0.00
P60867	40S ribosomal protein S20 OS=Mus musculus GN=Rps20 PE=1 SV=1	0.05	0.00	0.00	0.00
P61089	Ubiquitin-conjugating enzyme E2 N OS=Mus musculus GN=Ube2n PE=1 SV=1	0.02	0.00	0.02	0.00
P61168	D(2) dopamine receptor OS=Mus musculus GN=Drd2 PE=2 SV=2	0.04	0.03	1.00	1.00
P61255	60S ribosomal protein L26 OS=Mus musculus GN=Rpl26 PE=2 SV=1	0.00	0.00	0.05	1.00
P61358	60S ribosomal protein L27 OS=Mus musculus GN=Rpl27 PE=2 SV=2	1.00	0.04	0.02	0.02
P61750	ADP-ribosylation factor 4 OS=Mus musculus GN=Arf4 PE=2 SV=2	0.02	0.02	0.02	0.02
P61804	Dolichyl-diphosphooligosaccharide--protein glycosyltransferase subunit DAD1 OS=Mus musculus GN=Dad1 PE=	1.00	0.03	1.00	0.00
P61957	Small ubiquitin-related modifier 2 OS=Mus musculus GN=Sumo2 PE=2 SV=1	1.00	0.02	0.00	1.00
P61979	Heterogeneous nuclear ribonucleoprotein K OS=Mus musculus GN=Hnrnpk PE=1 SV=1	0.00	0.00	0.00	0.00
P61982	14-3-3 protein gamma OS=Mus musculus GN=Ywhag PE=1 SV=2	1.00	0.02	0.00	0.00
P62071	Ras-related protein R-Ras2 OS=Mus musculus GN=Rras2 PE=1 SV=1	1.00	1.00	1.00	0.02
P62082	40S ribosomal protein S7 OS=Mus musculus GN=Rps7 PE=2 SV=1	1.00	0.00	1.00	0.00
P62137	Serine/threonine-protein phosphatase PP1-alpha catalytic subunit OS=Mus musculus GN=Ppp1ca PE=1 SV=1	1.00	0.02	1.00	1.00
P62141	Serine/threonine-protein phosphatase PP1-beta catalytic subunit OS=Mus musculus GN=Ppp1cb PE=1 SV=3	1.00	0.02	1.00	1.00
P62204	Calmodulin OS=Mus musculus GN=Calm1 PE=1 SV=2	1.00	0.00	0.02	0.00
P62242	40S ribosomal protein S8 OS=Mus musculus GN=Rps8 PE=1 SV=2	0.02	0.00	0.00	0.00
P62245	40S ribosomal protein S15a OS=Mus musculus GN=Rps15a PE=2 SV=2	1.00	0.02	0.03	0.00
P62259	14-3-3 protein epsilon OS=Mus musculus GN=Ywhae PE=1 SV=1	0.00	0.00	0.00	0.00
P62264	40S ribosomal protein S14 OS=Mus musculus GN=Rps14 PE=2 SV=3	0.02	0.00	0.00	0.02
P62270	40S ribosomal protein S18 OS=Mus musculus GN=Rps18 PE=2 SV=3	0.00	0.00	0.00	0.07
P62281	40S ribosomal protein S11 OS=Mus musculus GN=Rps11 PE=2 SV=3	1.00	1.00	0.07	1.00
P62301	40S ribosomal protein S13 OS=Mus musculus GN=Rps13 PE=1 SV=2	1.00	0.05	1.00	1.00
P62305	Small nuclear ribonucleoprotein E OS=Mus musculus GN=Snrpe PE=2 SV=1	1.00	0.00	1.00	0.02
P62307	Small nuclear ribonucleoprotein F OS=Mus musculus GN=Snrpf PE=2 SV=1	1.00	0.00	0.02	1.00
P62309	Small nuclear ribonucleoprotein G OS=Mus musculus GN=Snrgp PE=1 SV=1	0.03	1.00	1.00	1.00
P62315	Small nuclear ribonucleoprotein Sm D1 OS=Mus musculus GN=Snrdp1 PE=2 SV=1	1.00	0.02	1.00	0.02
P62317	Small nuclear ribonucleoprotein Sm D2 OS=Mus musculus GN=Snrdp2 PE=2 SV=1	1.00	0.02	0.00	1.00
P62320	Small nuclear ribonucleoprotein Sm D3 OS=Mus musculus GN=Snrdp3 PE=1 SV=1	1.00	0.02	1.00	1.00
P62492	Ras-related protein Rab-11A OS=Mus musculus GN=Rab11a PE=1 SV=3	1.00	0.00	0.00	0.00
P62500	TSC22 domain family protein 1 OS=Mus musculus GN=Tsc22d1 PE=2 SV=2	1.00	0.02	0.02	1.00
P62702	40S ribosomal protein S4, X isoform OS=Mus musculus GN=Rps4x PE=2 SV=2	0.02	0.00	0.04	0.00
P62717	60S ribosomal protein L18a OS=Mus musculus GN=Rpl18a PE=1 SV=1	1.00	1.00	0.02	1.00
P62751	60S ribosomal protein L23a OS=Mus musculus GN=Rpl23a PE=1 SV=1	0.02	0.00	0.00	0.00
P62754	40S ribosomal protein S6 OS=Mus musculus GN=Rps6 PE=1 SV=1	1.00	0.00	1.00	1.00
P62806	Histone H4 OS=Mus musculus GN=Hist1h4a PE=1 SV=2	0.00	0.00	0.00	0.00
P62821	Ras-related protein Rab-1A OS=Mus musculus GN=Rab1A PE=1 SV=3	1.00	1.00	1.00	0.02
P62827	GTP-binding nuclear protein Ran OS=Mus musculus GN=Ran PE=1 SV=3	0.00	0.00	0.00	0.00
P62830	60S ribosomal protein L23 OS=Mus musculus GN=Rpl23 PE=1 SV=1	1.00	1.00	1.00	0.00
P62843	40S ribosomal protein S15 OS=Mus musculus GN=Rps15 PE=2 SV=2	1.00	0.02	0.02	0.05
P62849	40S ribosomal protein S24 OS=Mus musculus GN=Rps24 PE=1 SV=1	0.08	1.00	1.00	0.02

P62852	40S ribosomal protein S25 OS=Mus musculus GN=Rps25 PE=2 SV=1	1.00	0.00	0.02	0.00
P62855	40S ribosomal protein S26 OS=Mus musculus GN=Rps26 PE=2 SV=3	1.00	0.00	0.02	0.02
P62858	40S ribosomal protein S28 OS=Mus musculus GN=Rps28 PE=2 SV=1	1.00	0.00	0.02	0.02
P62869	Transcription elongation factor B polypeptide 2 OS=Mus musculus GN=Tceb2 PE=1 SV=1	1.00	0.02	1.00	1.00
P62880	Guanine nucleotide-binding protein G(I)/G(S)/G(T) subunit beta-2 OS=Mus musculus GN=Gnb2 PE=1 SV=3	1.00	0.05	1.00	0.02
P62889	60S ribosomal protein L30 OS=Mus musculus GN=Rpl30 PE=2 SV=2	1.00	0.05	1.00	0.00
P62900	60S ribosomal protein L31 OS=Mus musculus GN=Rpl31 PE=2 SV=1	0.07	0.02	0.02	0.03
P62908	40S ribosomal protein S3 OS=Mus musculus GN=Rps3 PE=1 SV=1	0.02	0.05	0.02	0.00
P62911	60S ribosomal protein L32 OS=Mus musculus GN=Rpl32 PE=2 SV=2	1.00	1.00	0.02	0.02
P62918	60S ribosomal protein L8 OS=Mus musculus GN=Rpl8 PE=2 SV=2	0.00	0.00	0.00	0.00
P62960	Nuclease-sensitive element-binding protein 1 OS=Mus musculus GN=Ybx1 PE=1 SV=3	1.00	0.00	0.02	0.00
P62962	Profilin-1 OS=Mus musculus GN=Pfn1 PE=1 SV=2	0.08	0.00	0.03	0.00
P62965	Cellular retinoic acid-binding protein 1 OS=Mus musculus GN=Crabp1 PE=1 SV=2	1.00	0.08	1.00	1.00
P62984	Ubiquitin-60S ribosomal protein L40 OS=Mus musculus GN=Uba52 PE=1 SV=2	0.00	0.00	0.00	0.00
P62996	Transformer-2 protein homolog beta OS=Mus musculus GN=Tra2b PE=1 SV=1	1.00	1.00	1.00	0.02
P63005	Platelet-activating factor acetylhydrolase IB subunit alpha OS=Mus musculus GN=Pafah1b1 PE=1 SV=2	1.00	1.00	0.02	1.00
P63017	Heat shock cognate 71 kDa protein OS=Mus musculus GN=Hspa8 PE=1 SV=1	0.00	0.00	0.00	0.00
P63028	Translationally-controlled tumor protein OS=Mus musculus GN=Tpt1 PE=1 SV=1	1.00	1.00	1.00	0.02
P63038	60 kDa heat shock protein, mitochondrial OS=Mus musculus GN=Hspd1 PE=1 SV=1	0.00	0.00	0.00	0.00
P63085	Mitogen-activated protein kinase 1 OS=Mus musculus GN=Mapk1 PE=1 SV=3	1.00	0.02	1.00	0.02
P63101	14-3-3 protein zeta/delta OS=Mus musculus GN=Ywhaz PE=1 SV=1	0.00	0.00	0.00	0.00
P63158	High mobility group protein B1 OS=Mus musculus GN=Hmgb1 PE=1 SV=2	0.00	0.00	0.00	0.00
P63163	Small nuclear ribonucleoprotein-associated protein N OS=Mus musculus GN=Snrpn PE=2 SV=1	0.05	1.00	1.00	1.00
P63242	Eukaryotic translation initiation factor 5A-1 OS=Mus musculus GN=Elf5a PE=1 SV=2	0.02	0.00	0.04	0.00
P63254	Cysteine-rich protein 1 OS=Mus musculus GN=Crip1 PE=2 SV=2	1.00	0.02	0.02	0.02
P63260	Actin, cytoplasmic 2 OS=Mus musculus GN=Actg1 PE=1 SV=1	0.00	0.00	0.00	0.00
P63276	40S ribosomal protein S17 OS=Mus musculus GN=Rps17 PE=1 SV=2	1.00	0.02	1.00	0.02
P63323	40S ribosomal protein S12 OS=Mus musculus GN=Rps12 PE=1 SV=2	1.00	0.02	0.02	0.03
P63325	40S ribosomal protein S10 OS=Mus musculus GN=Rps10 PE=1 SV=1	0.02	0.07	0.02	1.00
P67778	Prohibitin OS=Mus musculus GN=Phb PE=1 SV=1	1.00	1.00	0.02	1.00
P67871	Casein kinase II subunit beta OS=Mus musculus GN=Csnk2b PE=1 SV=1	1.00	1.00	0.06	1.00
P67984	60S ribosomal protein L22 OS=Mus musculus GN=Rpl22 PE=2 SV=2	1.00	0.00	0.00	0.00
P68037	Ubiquitin-conjugating enzyme E2 L3 OS=Mus musculus GN=Ube2l3 PE=2 SV=1	1.00	0.04	0.06	1.00
P68040	Guanine nucleotide-binding protein subunit beta-2-like 1 OS=Mus musculus GN=Gnb2l1 PE=1 SV=3	1.00	1.00	0.02	0.05
P68134	Actin, alpha skeletal muscle OS=Mus musculus GN=Acta1 PE=1 SV=1	0.00	0.00	0.00	0.00
P68254	14-3-3 protein theta OS=Mus musculus GN=Ywhaq PE=1 SV=1	0.00	0.00	0.02	0.00
P68369	Tubulin alpha-1A chain OS=Mus musculus GN=Tuba1a PE=1 SV=1	0.07	0.00	0.02	0.00
P68372	Tubulin beta-2C chain OS=Mus musculus GN=Tubb2c PE=1 SV=1	1.00	1.00	1.00	0.00
P68433	Histone H3.1 OS=Mus musculus GN=Hist1h3a PE=1 SV=2	0.00	0.00	0.03	0.02
P68510	14-3-3 protein eta OS=Mus musculus GN=Ywhah PE=1 SV=2	0.03	0.00	0.00	0.00
P70121	Zinc fingers and homeoboxes protein 1 OS=Mus musculus GN=Zhx1 PE=1 SV=2	1.00	0.07	1.00	1.00
P70168	Importin subunit beta-1 OS=Mus musculus GN=Kpnb1 PE=1 SV=2	1.00	1.00	0.04	0.00
P70195	Proteasome subunit beta type-7 OS=Mus musculus GN=Psb7 PE=1 SV=1	1.00	0.00	0.03	1.00
P70296	Phosphatidylethanolamine-binding protein 1 OS=Mus musculus GN=Pebp1 PE=1 SV=3	1.00	0.04	1.00	0.00
P70333	Heterogeneous nuclear ribonucleoprotein H2 OS=Mus musculus GN=Hnrnp2 PE=1 SV=1	1.00	0.00	0.02	0.00
P70349	Histidine triad nucleotide-binding protein 1 OS=Mus musculus GN=Hint1 PE=1 SV=3	1.00	1.00	0.02	0.02
P70372	ELAV-like protein 1 OS=Mus musculus GN=Elavl1 PE=1 SV=2	1.00	0.02	0.02	1.00
P70441	Na(+)/H(+) exchange regulatory cofactor NHE-RF1 OS=Mus musculus GN=Slc9a3r1 PE=1 SV=3	1.00	1.00	0.02	0.00
P80313	T-complex protein 1 subunit eta OS=Mus musculus GN=Cct7 PE=1 SV=1	1.00	1.00	0.02	1.00
P80315	T-complex protein 1 subunit delta OS=Mus musculus GN=Cct4 PE=1 SV=3	1.00	0.06	1.00	0.04
P80316	T-complex protein 1 subunit epsilon OS=Mus musculus GN=Cct5 PE=1 SV=1	1.00	1.00	1.00	0.02
P83917	Chromobox protein homolog 1 OS=Mus musculus GN=Cbx1 PE=1 SV=1	1.00	0.02	0.04	0.02
P84078	ADP-ribosylation factor 1 OS=Mus musculus GN=Arf1 PE=1 SV=2	1.00	0.00	0.04	0.00
P84099	60S ribosomal protein L19 OS=Mus musculus GN=Rpl19 PE=1 SV=1	0.02	0.02	0.02	0.00
P84104	Serine/arginine-rich splicing factor 3 OS=Mus musculus GN=Srsf3 PE=2 SV=1	0.02	0.02	1.00	0.02
P84228	Histone H3.2 OS=Mus musculus GN=Hist1h3b PE=1 SV=2	0.02	0.02	1.00	0.02
P84244	Histone H3.3 OS=Mus musculus GN=H3f3a PE=1 SV=2	0.02	0.02	1.00	0.02
P97300	Neuroplastin OS=Mus musculus GN=Nptn PE=1 SV=3	1.00	1.00	0.07	0.08
P97310	DNA replication licensing factor MCM2 OS=Mus musculus GN=Mcm2 PE=1 SV=3	1.00	1.00	0.02	1.00
P97315	Cysteine and glycine-rich protein 1 OS=Mus musculus GN=Csrp1 PE=1 SV=3	1.00	1.00	0.05	1.00
P97350	Plakophilin-1 OS=Mus musculus GN=Pkp1 PE=1 SV=1	1.00	1.00	1.00	0.02
P97351	40S ribosomal protein S3a OS=Mus musculus GN=Rps3a PE=1 SV=3	1.00	0.00	0.00	0.02
P97352	Protein S100-A13 OS=Mus musculus GN=S100a13 PE=1 SV=1	1.00	1.00	0.02	0.02
P97371	Proteasome activator complex subunit 1 OS=Mus musculus GN=Psmc1 PE=2 SV=2	0.03	1.00	0.02	0.02
P97372	Proteasome activator complex subunit 2 OS=Mus musculus GN=Psmc2 PE=2 SV=4	1.00	1.00	1.00	0.05
P97429	Annexin A4 OS=Mus musculus GN=Anxa4 PE=2 SV=4	1.00	0.04	0.00	0.00
P97447	Four and a half LIM domains protein 1 OS=Mus musculus GN=Fhl1 PE=2 SV=3	1.00	1.00	0.02	0.02
P97449	Aminopeptidase N OS=Mus musculus GN=Anpep PE=1 SV=4	1.00	1.00	0.03	0.00
P97450	ATP synthase-coupling factor 6, mitochondrial OS=Mus musculus GN=Atp5j PE=1 SV=1	1.00	0.04	1.00	1.00
P97497	Sex hormone-binding globulin OS=Mus musculus GN=Shbg PE=2 SV=1	1.00	0.07	1.00	1.00
P97807	Fumarate hydratase, mitochondrial OS=Mus musculus GN=Fh PE=1 SV=3	1.00	0.02	1.00	0.00

P97855	Ras GTPase-activating protein-binding protein 1 OS=Mus musculus GN=G3bp1 PE=1 SV=1	1.00	1.00	1.00	0.00
P99024	Tubulin beta-5 chain OS=Mus musculus GN=Tubb5 PE=1 SV=1	1.00	0.02	0.00	0.00
P99026	Proteasome subunit beta type-4 OS=Mus musculus GN=Psb4 PE=1 SV=1	1.00	0.00	1.00	1.00
P99027	60S acidic ribosomal protein P2 OS=Mus musculus GN=Rplp2 PE=1 SV=3	0.00	0.00	0.02	0.00
P99029	Peroxisomal protein OS=Mus musculus GN=Prdx5 PE=1 SV=2	1.00	0.02	1.00	0.00
Q00519	Xanthine dehydrogenase/oxidase OS=Mus musculus GN=Xdh PE=1 SV=5	1.00	1.00	1.00	0.02
Q00915	Retinol-binding protein 1 OS=Mus musculus GN=Rbp1 PE=2 SV=2	1.00	0.02	0.02	0.00
Q00P19	Heterogeneous nuclear ribonucleoprotein U-like protein 2 OS=Mus musculus GN=Hnrnpul2 PE=1 SV=1	1.00	1.00	0.02	1.00
Q01730	Ras suppressor protein 1 OS=Mus musculus GN=Rsu1 PE=2 SV=3	1.00	1.00	1.00	0.02
Q01768	Nucleoside diphosphate kinase B OS=Mus musculus GN=Nme2 PE=1 SV=1	0.02	0.00	0.00	0.00
Q01853	Transitional endoplasmic reticulum ATPase OS=Mus musculus GN=Vcp PE=1 SV=4	0.03	0.00	1.00	0.00
Q02053	Ubiquitin-like modifier-activating enzyme 1 OS=Mus musculus GN=Uba1 PE=1 SV=1	0.08	0.00	0.00	0.00
Q02257	Junction plakoglobin OS=Mus musculus GN=Jup PE=1 SV=3	1.00	1.00	1.00	0.00
Q03265	ATP synthase subunit alpha, mitochondrial OS=Mus musculus GN=Atp5a1 PE=1 SV=1	0.07	0.00	0.02	0.00
Q04447	Creatine kinase B-type OS=Mus musculus GN=Ckb PE=1 SV=1	1.00	1.00	1.00	0.00
Q04857	Collagen alpha-1(VI) chain OS=Mus musculus GN=Col6a1 PE=2 SV=1	1.00	1.00	0.02	0.04
Q05144	Ras-related C3 botulinum toxin substrate 2 OS=Mus musculus GN=Rac2 PE=2 SV=1	1.00	1.00	0.02	0.02
Q05186	Reticulocalbin-1 OS=Mus musculus GN=Rcn1 PE=1 SV=1	0.02	0.00	0.02	0.00
Q05816	Fatty acid-binding protein, epidermal OS=Mus musculus GN=Fabp5 PE=1 SV=3	0.02	0.00	0.00	0.00
Q06185	ATP synthase subunit e, mitochondrial OS=Mus musculus GN=Atp5i PE=1 SV=2	1.00	0.02	0.02	1.00
Q07113	Cation-independent mannose-6-phosphate receptor OS=Mus musculus GN=Igf2r PE=1 SV=1	0.02	0.00	0.00	1.00
Q07475	Neuronal protein 3.1 OS=Mus musculus GN=P311 PE=2 SV=1	1.00	1.00	0.05	1.00
Q08189	Protein-glutamine gamma-glutamyltransferase E OS=Mus musculus GN=Tgm3 PE=1 SV=2	1.00	1.00	1.00	0.03
Q09163	Protein delta homolog 1 OS=Mus musculus GN=Dk1 PE=1 SV=1	1.00	0.04	1.00	1.00
Q0VBK2	Keratin, type II cytoskeletal 80 OS=Mus musculus GN=Krt80 PE=2 SV=1	1.00	1.00	1.00	0.00
Q2VIS4	Filaggrin-2 OS=Mus musculus GN=Flg2 PE=1 SV=2	1.00	1.00	1.00	0.02
Q3TEA8	Heterochromatin protein 1-binding protein 3 OS=Mus musculus GN=Hp1bp3 PE=1 SV=1	1.00	1.00	1.00	0.04
Q3THE2	Myosin regulatory light chain 12B OS=Mus musculus GN=Myl12b PE=1 SV=2	1.00	0.00	0.00	0.00
Q3THW5	Histone H2A.V OS=Mus musculus GN=H2afv PE=1 SV=3	0.02	0.05	1.00	0.00
Q3TJ91	Lethal(2) giant larvae protein homolog 2 OS=Mus musculus GN=Llg2 PE=2 SV=2	1.00	0.04	1.00	1.00
Q3TML0	Uncharacterized protein OS=Mus musculus GN=Pdia6 PE=2 SV=1	0.02	0.00	0.00	0.00
Q3TWW8	Splicing factor, arginine/serine-rich 6 OS=Mus musculus GN=Srsf6 PE=2 SV=1	1.00	0.02	0.02	1.00
Q3TYA6	RIKEN cDNA 4930548G07 OS=Mus musculus GN=Mphosph8 PE=2 SV=1	0.04	1.00	1.00	1.00
Q3U0V1	Far upstream element-binding protein 2 OS=Mus musculus GN=Khsrp PE=1 SV=2	1.00	0.00	0.00	0.03
Q3U1J4	DNA damage-binding protein 1 OS=Mus musculus GN=Ddb1 PE=1 SV=2	1.00	1.00	0.08	1.00
Q3U7R1	Extended synaptotagmin-1 OS=Mus musculus GN=Esy1 PE=2 SV=2	1.00	1.00	1.00	0.03
Q3UHD9	Arf-GAP with GTPase, ANK repeat and PH domain-containing protein 2 OS=Mus musculus GN=Agap2 PE=1 SV=1	1.00	0.04	1.00	1.00
Q3UHX2	28 kDa heat- and acid-stable phosphoprotein OS=Mus musculus GN=Pdap1 PE=1 SV=1	1.00	1.00	1.00	0.05
Q3UJB0	Splicing factor 3b, subunit 2 OS=Mus musculus GN=Sf3b2 PE=2 SV=1	0.04	1.00	1.00	1.00
Q3USB7	Inactive phospholipase C-like protein 1 OS=Mus musculus GN=Plcl1 PE=1 SV=3	1.00	0.06	1.00	1.00
Q3UW53	Protein Niban OS=Mus musculus GN=Fam129a PE=1 SV=2	1.00	1.00	1.00	0.00
Q3V0P4	Gene model 1661 OS=Mus musculus GN=Gm1661 PE=2 SV=1	1.00	1.00	0.08	1.00
Q3V0Q1	Dynein heavy chain 12, axonemal OS=Mus musculus GN=Dnah12 PE=1 SV=2	1.00	1.00	1.00	0.05
Q3V1T4	Prolyl 3-hydroxylase 1 OS=Mus musculus GN=Lepre1 PE=2 SV=2	1.00	1.00	1.00	0.02
Q4KL66	Pregnancy-specific glycoprotein 28 OS=Mus musculus GN=Psg28 PE=2 SV=1	0.01	1.00	1.00	1.00
Q4U4S6	Xin actin-binding repeat-containing protein 2 OS=Mus musculus GN=Xirp2 PE=1 SV=1	1.00	1.00	1.00	0.07
Q4V9X0	Ppp3ca protein OS=Mus musculus GN=Ppp3ca PE=2 SV=1	1.00	0.07	1.00	1.00
Q501J6	Probable ATP-dependent RNA helicase DDX17 OS=Mus musculus GN=Ddx17 PE=2 SV=1	0.02	0.02	1.00	0.00
Q504N0	Carboxypeptidase A2 OS=Mus musculus GN=Cpa2 PE=2 SV=1	1.00	0.00	0.03	1.00
Q52KH6	Novel protein OS=Mus musculus GN=DXBay18 PE=2 SV=1	1.00	1.00	1.00	0.03
Q58E70	Tpm3 protein OS=Mus musculus GN=Tpm3 PE=2 SV=1	1.00	0.00	0.00	0.00
Q5BKQ4	Pancreatic lipase-related protein 1 OS=Mus musculus GN=Pnlipr1 PE=2 SV=2	0.02	0.00	0.00	0.00
Q5DTT4	Retrotransposon gag domain-containing protein 4 OS=Mus musculus GN=Rgag4 PE=2 SV=2	1.00	0.01	1.00	1.00
Q5F4T0	Transient receptor potential cation channel, subfamily M, member 3 OS=Mus musculus GN=Trpm3 PE=2 SV=1	1.00	0.02	1.00	1.00
Q5XJY5	Coatomer subunit delta OS=Mus musculus GN=Arcn1 PE=2 SV=2	1.00	1.00	1.00	0.02
Q60605	Myosin light polypeptide 6 OS=Mus musculus GN=Myl6 PE=1 SV=3	1.00	0.02	0.00	0.00
Q60662	A-kinase anchor protein 4 OS=Mus musculus GN=Akap4 PE=1 SV=1	0.06	0.02	1.00	0.04
Q60668	Heterogeneous nuclear ribonucleoprotein D0 OS=Mus musculus GN=Hnrnpd PE=1 SV=2	1.00	0.00	0.02	0.00
Q60692	Proteasome subunit beta type-6 OS=Mus musculus GN=Psb6 PE=1 SV=3	1.00	0.00	0.02	0.02
Q60715	Prolyl 4-hydroxylase subunit alpha-1 OS=Mus musculus GN=P4ha1 PE=2 SV=2	1.00	1.00	1.00	0.00
Q60716	Prolyl 4-hydroxylase subunit alpha-2 OS=Mus musculus GN=P4ha2 PE=2 SV=1	1.00	0.02	0.02	1.00
Q60739	BAG family molecular chaperone regulator 1 OS=Mus musculus GN=Bag1 PE=1 SV=3	1.00	0.07	0.02	0.03
Q60751	Insulin-like growth factor 1 receptor OS=Mus musculus GN=Igf1r PE=1 SV=3	1.00	0.01	1.00	1.00
Q60817	Nascent polypeptide-associated complex subunit alpha OS=Mus musculus GN=Naca PE=1 SV=1	1.00	0.00	1.00	0.02
Q60854	Serpin B6 OS=Mus musculus GN=Serpnb6 PE=2 SV=1	1.00	1.00	1.00	0.00
Q60864	Stress-induced-phosphoprotein 1 OS=Mus musculus GN=Stip1 PE=1 SV=1	1.00	0.02	0.04	1.00
Q60865	Caprin-1 OS=Mus musculus GN=Caprin1 PE=1 SV=2	1.00	1.00	1.00	0.07
Q60870	Receptor expression-enhancing protein 5 OS=Mus musculus GN=Reep5 PE=1 SV=1	1.00	1.00	0.02	1.00
Q60931	Voltage-dependent anion-selective channel protein 3 OS=Mus musculus GN=Vdac3 PE=1 SV=1	1.00	1.00	1.00	0.07
Q60932	Voltage-dependent anion-selective channel protein 1 OS=Mus musculus GN=Vdac1 PE=1 SV=3	1.00	0.00	0.00	0.00
Q60997	Deleted in malignant brain tumors 1 protein OS=Mus musculus GN=Dmbt1 PE=1 SV=2	1.00	1.00	0.00	1.00

Q61029	Lamina-associated polypeptide 2, isoforms beta/delta/epsilon/gamma OS=Mus musculus GN=Tmpo PE=1 SV=4	1.00	0.02	0.00	0.00
Q61035	Histidyl-tRNA synthetase, cytoplasmic OS=Mus musculus GN=Hars PE=2 SV=2	1.00	0.02	0.02	1.00
Q61171	Peroxiredoxin-2 OS=Mus musculus GN=Prdx2 PE=1 SV=3	0.02	0.00	0.02	0.00
Q61207	Sulfated glycoprotein 1 OS=Mus musculus GN=Psap PE=1 SV=2	1.00	0.02	0.08	1.00
Q61409	cGMP-inhibited 3',5'-cyclic phosphodiesterase B OS=Mus musculus GN=Pde3b PE=1 SV=2	1.00	0.02	1.00	1.00
Q61425	Hydroxyacyl-coenzyme A dehydrogenase, mitochondrial OS=Mus musculus GN=Hadh PE=1 SV=2	1.00	0.00	1.00	1.00
Q61545	RNA-binding protein EWS OS=Mus musculus GN=Ewsr1 PE=1 SV=2	1.00	0.00	1.00	1.00
Q61553	Fascin OS=Mus musculus GN=Fscn1 PE=1 SV=4	1.00	0.00	0.00	0.00
Q61598	Rab GDP dissociation inhibitor beta OS=Mus musculus GN=Gdi2 PE=1 SV=1	1.00	1.00	1.00	0.02
Q61599	Rho GDP-dissociation inhibitor 2 OS=Mus musculus GN=Arhgdib PE=1 SV=3	1.00	1.00	1.00	0.02
Q61753	D-3-phosphoglycerate dehydrogenase OS=Mus musculus GN=Phgdh PE=1 SV=3	1.00	0.03	0.00	1.00
Q61792	LIM and SH3 domain protein 1 OS=Mus musculus GN=Lasp1 PE=1 SV=1	1.00	1.00	1.00	0.02
Q61879	Myosin-10 OS=Mus musculus GN=Myh10 PE=1 SV=2	1.00	1.00	1.00	0.02
Q61937	Nucleophosmin OS=Mus musculus GN=Npm1 PE=1 SV=1	0.00	0.00	0.00	0.00
Q61990	Poly(rC)-binding protein 2 OS=Mus musculus GN=Pcbp2 PE=1 SV=1	1.00	1.00	1.00	0.00
Q62048	Astrocytic phosphoprotein PEA-15 OS=Mus musculus GN=Pea15 PE=1 SV=1	1.00	0.02	0.02	0.02
Q62093	Serine/arginine-rich splicing factor 2 OS=Mus musculus GN=Srsf2 PE=1 SV=4	1.00	0.02	1.00	0.02
Q62148	Retinal dehydrogenase 2 OS=Mus musculus GN=Aldh1a2 PE=1 SV=2	1.00	0.03	1.00	0.02
Q62167	ATP-dependent RNA helicase DDX3X OS=Mus musculus GN=Ddx3x PE=1 SV=3	0.08	1.00	0.05	1.00
Q62188	Dihydropyrimidinase-related protein 3 OS=Mus musculus GN=Dpysl3 PE=1 SV=1	1.00	0.03	0.02	0.00
Q62189	U1 small nuclear ribonucleoprotein A OS=Mus musculus GN=Snrpa PE=2 SV=3	1.00	0.02	0.03	1.00
Q62261	Spectrin beta chain, brain 1 OS=Mus musculus GN=Sptbn1 PE=1 SV=2	1.00	0.05	0.00	0.00
Q62318	Transcription intermediary factor 1-beta OS=Mus musculus GN=Trim28 PE=1 SV=3	1.00	0.03	1.00	1.00
Q62376	U1 small nuclear ribonucleoprotein 70 kDa OS=Mus musculus GN=Snrnp70 PE=1 SV=2	1.00	0.08	1.00	1.00
Q62446	Peptidyl-prolyl cis-trans isomerase FKBP3 OS=Mus musculus GN=Fkbp3 PE=1 SV=2	1.00	0.02	1.00	1.00
Q62465	Synaptic vesicle membrane protein VAT-1 homolog OS=Mus musculus GN=Vat1 PE=1 SV=3	1.00	0.02	0.00	0.00
Q63836	Selenium-binding protein 2 OS=Mus musculus GN=Selenbp2 PE=1 SV=2	1.00	0.02	0.02	0.02
Q63918	Serum deprivation-response protein OS=Mus musculus GN=Sdpr PE=1 SV=3	1.00	1.00	1.00	0.03
Q64012	RNA-binding protein Raly OS=Mus musculus GN=Raly PE=1 SV=3	0.04	1.00	0.07	1.00
Q640L3	Cell cycle progression protein 1 OS=Mus musculus GN=Ccpg1 PE=1 SV=2	1.00	0.04	1.00	1.00
Q64285	Bile salt-activated lipase OS=Mus musculus GN=Cel PE=1 SV=1	1.00	0.00	0.02	0.00
Q64314	Hematopoietic progenitor cell antigen CD34 OS=Mus musculus GN=Cd34 PE=1 SV=1	1.00	0.07	1.00	1.00
Q64433	10 kDa heat shock protein, mitochondrial OS=Mus musculus GN=Hspe1 PE=1 SV=2	0.02	0.00	0.00	0.00
Q64522	Histone H2A type 2-B OS=Mus musculus GN=Hist2h2ab PE=1 SV=3	0.00	0.00	1.00	0.00
Q64727	Vinculin OS=Mus musculus GN=Vcl PE=1 SV=4	1.00	0.02	0.00	0.00
Q68FD5	Clathrin heavy chain 1 OS=Mus musculus GN=Cltc PE=1 SV=3	1.00	1.00	1.00	0.00
Q6A0C2	MKIAA0136 protein (Fragment) OS=Mus musculus GN=Morc3 PE=2 SV=1	1.00	0.05	1.00	1.00
Q6B966	NACHT, LRR and PYD domains-containing protein 14 OS=Mus musculus GN=Nlrp14 PE=2 SV=2	1.00	1.00	1.00	0.03
Q6IE03	Sentrin/SUMO-specific protease 15 OS=Mus musculus GN=Gm9839 PE=2 SV=1	0.07	0.08	1.00	0.07
Q6IFZ6	Keratin, type II cytoskeletal 1b OS=Mus musculus GN=Krt77 PE=1 SV=1	0.08	1.00	1.00	0.00
Q6IRU2	Tropomyosin alpha-4 chain OS=Mus musculus GN=Tpm4 PE=2 SV=3	1.00	0.00	0.00	0.00
Q6P069	Sorcini OS=Mus musculus GN=Sri PE=1 SV=1	1.00	1.00	0.04	1.00
Q6P5G0	Mitogen-activated protein kinase 4 OS=Mus musculus GN=Mapk4 PE=2 SV=1	1.00	1.00	1.00	0.08
Q6P9N8	Trafficking protein, kinesin binding 2 OS=Mus musculus GN=Trak2 PE=2 SV=1	1.00	0.03	1.00	0.05
Q6PDG5	SWI/SNF complex subunit SMARCC2 OS=Mus musculus GN=Smarcc2 PE=1 SV=2	1.00	0.08	1.00	1.00
Q6PDM2	Serine/arginine-rich splicing factor 1 OS=Mus musculus GN=Srsf1 PE=1 SV=3	0.00	0.00	0.02	1.00
Q6PDN3	Myosin light chain kinase, smooth muscle OS=Mus musculus GN=Mylk PE=1 SV=3	1.00	1.00	0.08	1.00
Q6PGA0	REST corepressor 3 OS=Mus musculus GN=Rcor3 PE=2 SV=2	1.00	1.00	0.08	1.00
Q6PHZ5	Putative RNA-binding protein 15B OS=Mus musculus GN=Rbm15b PE=1 SV=2	1.00	1.00	0.07	1.00
Q6PJ18	Tpm2 protein OS=Mus musculus GN=Tpm2 PE=2 SV=1	1.00	0.02	1.00	0.02
Q6WKZ8	E3 ubiquitin-protein ligase UBR2 OS=Mus musculus GN=Ubr2 PE=1 SV=2	1.00	0.01	1.00	0.01
Q6VVG3	BTB/POZ domain-containing protein KCTD12 OS=Mus musculus GN=Kctd12 PE=1 SV=1	1.00	0.00	0.02	0.02
Q6ZPY7	Lysine-specific demethylase 3B OS=Mus musculus GN=Kdm3b PE=1 SV=2	0.04	0.05	1.00	1.00
Q6ZWN5	40S ribosomal protein S9 OS=Mus musculus GN=Rps9 PE=2 SV=3	1.00	0.02	0.02	0.03
Q6ZWU9	40S ribosomal protein S27 OS=Mus musculus GN=Rps27 PE=1 SV=3	1.00	1.00	0.02	1.00
Q6ZWV3	60S ribosomal protein L10 OS=Mus musculus GN=Rpl10 PE=2 SV=3	1.00	1.00	1.00	0.00
Q6ZWV7	60S ribosomal protein L35 OS=Mus musculus GN=Rpl35 PE=2 SV=1	0.02	0.02	0.02	0.02
Q6ZWX6	Eukaryotic translation initiation factor 2 subunit 1 OS=Mus musculus GN=Eif2s1 PE=1 SV=3	1.00	1.00	0.02	1.00
Q78PY7	Staphylococcal nuclease domain-containing protein 1 OS=Mus musculus GN=Snd1 PE=1 SV=1	1.00	0.00	0.00	1.00
Q792Y9	MCG140783 OS=Mus musculus GN=Gm5771 PE=2 SV=1	1.00	0.02	0.02	1.00
Q792Z0	Trypsinogen 11 OS=Mus musculus GN=Prss3 PE=2 SV=1	1.00	0.00	0.00	1.00
Q792Z1	MCG140784 OS=Mus musculus GN=Try10 PE=2 SV=1	1.00	0.00	0.02	0.02
Q77MK9	Heterogeneous nuclear ribonucleoprotein Q OS=Mus musculus GN=Synrcp PE=1 SV=2	1.00	0.02	0.00	0.00
Q7TNC4	Putative RNA-binding protein Luc7-like 2 OS=Mus musculus GN=Luc7l2 PE=1 SV=1	1.00	0.02	1.00	1.00
Q7TPR4	Alpha-actinin-1 OS=Mus musculus GN=Actn1 PE=1 SV=1	0.03	1.00	1.00	0.00
Q7TPZ8	Carboxypeptidase A1 OS=Mus musculus GN=Cpa1 PE=2 SV=1	0.02	0.00	0.00	0.00
Q80TN7	Neuron navigator 3 OS=Mus musculus GN=Nav3 PE=1 SV=2	0.04	1.00	1.00	1.00
Q80VM4	Zinc finger protein 579 OS=Mus musculus GN=Znf579 PE=1 SV=1	1.00	1.00	1.00	0.08
Q80WJ7	Protein LYRIC OS=Mus musculus GN=Mtdh PE=1 SV=1	1.00	1.00	0.05	1.00
Q80WT0	Junctophilin-4 OS=Mus musculus GN=Jph4 PE=2 SV=1	1.00	0.07	0.07	1.00
Q80X90	Filamin-B OS=Mus musculus GN=Finb PE=1 SV=3	1.00	1.00	0.07	0.00

Q80XD0	BC051142 protein OS=Mus musculus GN=BC051142 PE=2 SV=1	1.00	0.01	1.00	1.00
Q80XU3	Nuclear ubiquitous casein and cyclin-dependent kinases substrate OS=Mus musculus GN=Nucks1 PE=1 SV=1	1.00	0.02	1.00	1.00
Q8BFR5	Elongation factor Tu, mitochondrial OS=Mus musculus GN=Tufm PE=1 SV=1	1.00	1.00	0.03	1.00
Q8BFS6	Calcineurin-like phosphoesterase domain-containing protein 1 OS=Mus musculus GN=Cpped1 PE=2 SV=1	1.00	1.00	0.05	1.00
Q8BG05	Heterogeneous nuclear ribonucleoprotein A3 OS=Mus musculus GN=Hnrmpa3 PE=1 SV=1	0.00	0.00	0.00	0.00
Q8BGD9	Eukaryotic translation initiation factor 4B OS=Mus musculus GN=EIF4B PE=1 SV=1	0.04	1.00	1.00	1.00
Q8BGS0	Protein MAK16 homolog OS=Mus musculus GN=Mak16 PE=2 SV=1	1.00	1.00	1.00	0.07
Q8BH64	EH domain-containing protein 2 OS=Mus musculus GN=Ehd2 PE=1 SV=1	1.00	1.00	1.00	0.00
Q8BH95	Enoyl-CoA hydratase, mitochondrial OS=Mus musculus GN=Echs1 PE=1 SV=1	1.00	1.00	0.02	1.00
Q8BH97	Reticulocalbin-3 OS=Mus musculus GN=Rcn3 PE=2 SV=1	1.00	0.00	0.02	0.00
Q8BHN3	Neutral alpha-glucosidase AB OS=Mus musculus GN=Ganab PE=1 SV=1	1.00	0.04	1.00	1.00
Q8BJU0	Small glutamine-rich tetrapeptide repeat-containing protein alpha OS=Mus musculus GN=Sgta PE=1 SV=2	1.00	0.07	1.00	1.00
Q8BK67	Protein RCC2 OS=Mus musculus GN=Rcc2 PE=2 SV=1	1.00	0.06	1.00	1.00
Q8BL97	Serine/arginine-rich splicing factor 7 OS=Mus musculus GN=Srsf7 PE=1 SV=1	1.00	0.00	0.03	1.00
Q8BМК4	Cytoskeleton-associated protein 4 OS=Mus musculus GN=Ckap4 PE=2 SV=2	0.00	0.00	0.00	0.00
Q8BMS1	Trifunctional enzyme subunit alpha, mitochondrial OS=Mus musculus GN=Hadha PE=1 SV=1	1.00	1.00	0.00	0.00
Q8BP67	60S ribosomal protein L24 OS=Mus musculus GN=Rpl24 PE=2 SV=2	1.00	1.00	0.07	0.02
Q8BP92	Reticulocalbin-2 OS=Mus musculus GN=Rcn2 PE=2 SV=1	1.00	1.00	1.00	0.02
Q8BSY0	Aspartyl/asparaginyl beta-hydroxylase OS=Mus musculus GN=Asph PE=2 SV=1	1.00	1.00	1.00	0.02
Q8BTF7	DNA ligase 4 OS=Mus musculus GN=Lig4 PE=2 SV=1	1.00	1.00	0.07	1.00
Q8BTM8	Filamin-A OS=Mus musculus GN=Flna PE=1 SV=5	1.00	0.00	0.00	0.00
Q8BUE4	Apoptosis-inducing factor 2 OS=Mus musculus GN=Aifm2 PE=2 SV=1	1.00	1.00	0.05	0.02
Q8BWT1	3-ketoacyl-CoA thiolase, mitochondrial OS=Mus musculus GN=Acaa2 PE=1 SV=3	1.00	1.00	1.00	0.00
Q8BX57	PX domain-containing protein kinase-like protein OS=Mus musculus GN=Pxx PE=1 SV=2	1.00	0.08	1.00	1.00
Q8C0M9	L-asparaginase OS=Mus musculus GN=Asrgl1 PE=1 SV=1	1.00	1.00	1.00	0.04
Q8C138	Uncharacterized protein C6orf105 homolog OS=Mus musculus PE=2 SV=2	1.00	1.00	1.00	0.03
Q8C1B7	Septin-11 OS=Mus musculus GN=Sept11 PE=1 SV=4	1.00	0.00	0.02	0.02
Q8C635	Glycerol kinase-like 1 OS=Mus musculus GN=Gykl1 PE=2 SV=1	1.00	1.00	0.07	1.00
Q8CCT4	Transcription elongation factor A protein-like 5 OS=Mus musculus GN=Tcea5 PE=1 SV=1	0.02	1.00	1.00	1.00
Q8CDN6	Thioredoxin-like protein 1 OS=Mus musculus GN=Txn1f1 PE=1 SV=3	1.00	0.02	0.00	1.00
Q8CE23	Orexigenic neuropeptide QRFP OS=Mus musculus GN=Qrfp PE=2 SV=1	1.00	0.08	1.00	1.00
Q8CGP2	Histone H2B type 1-P OS=Mus musculus GN=Hist1h2bp PE=1 SV=3	0.00	0.00	0.00	0.00
Q8CIE6	Coatamer subunit alpha OS=Mus musculus GN=Copa PE=1 SV=2	1.00	1.00	1.00	0.02
Q8K0E8	Fibrinogen beta chain OS=Mus musculus GN=Fgb PE=2 SV=1	1.00	1.00	1.00	0.02
Q8K0Z5	Tropomyosin 3, gamma OS=Mus musculus GN=Tpm3 PE=2 SV=1	1.00	0.03	0.05	0.08
Q8K2B3	Succinate dehydrogenase [ubiquinone] flavoprotein subunit, mitochondrial OS=Mus musculus GN=Sdha PE=1 S	0.03	0.02	0.02	0.03
Q8K3P1	P2X purinoceptor OS=Mus musculus GN=P2rx2 PE=2 SV=2	0.05	0.08	1.00	1.00
Q8K4L4	Protein POF1B OS=Mus musculus GN=Pof1b PE=1 SV=3	1.00	1.00	0.02	1.00
Q8K4Z0	Leucine-rich repeat LGI family member 2 OS=Mus musculus GN=Lgi2 PE=2 SV=1	1.00	1.00	1.00	0.01
Q8QZT1	Acetyl-CoA acetyltransferase, mitochondrial OS=Mus musculus GN=Acat1 PE=1 SV=1	1.00	0.00	1.00	0.00
Q8R050	Eukaryotic peptide chain release factor GTP-binding subunit ERF3A OS=Mus musculus GN=Gspst1 PE=1 SV=1	1.00	0.07	1.00	1.00
Q8R081	Heterogeneous nuclear ribonucleoprotein L OS=Mus musculus GN=Hnrpl PE=1 SV=2	1.00	0.00	1.00	1.00
Q8R1M2	Histone H2A.J OS=Mus musculus GN=H2afj PE=2 SV=1	0.02	0.02	1.00	0.02
Q8R207	Small subunit of serine palmitoyltransferase A OS=Mus musculus GN=Ssspta PE=2 SV=2	1.00	0.02	0.02	0.02
Q8R370	Usher syndrome type-1C protein-binding protein 1 OS=Mus musculus GN=Ushbp1 PE=1 SV=2	1.00	1.00	0.03	1.00
Q8R5C5	Beta-centractin OS=Mus musculus GN=Actr1b PE=1 SV=1	1.00	0.02	0.03	1.00
Q8VCQ6	Phosphatidylcholine:ceramide cholinephosphotransferase 1 OS=Mus musculus GN=Sgms1 PE=1 SV=2	0.02	1.00	1.00	1.00
Q8VCQ8	Caldesmon 1 OS=Mus musculus GN=Cald1 PE=2 SV=1	1.00	0.00	0.02	0.00
Q8VDD5	Myosin-9 OS=Mus musculus GN=Myh9 PE=1 SV=4	1.00	1.00	0.00	0.00
Q8VDN2	Sodium/potassium-transporting ATPase subunit alpha-1 OS=Mus musculus GN=Atp1a1 PE=1 SV=1	1.00	1.00	1.00	0.02
Q8VDP4	Protein KIAA1967 homolog OS=Mus musculus PE=1 SV=2	1.00	1.00	1.00	0.02
Q8VE97	Serine/arginine-rich splicing factor 4 OS=Mus musculus GN=Srsf4 PE=2 SV=1	0.07	0.03	0.02	0.02
Q8VEK3	Heterogeneous nuclear ribonucleoprotein U OS=Mus musculus GN=Hnrmpu PE=1 SV=1	0.03	0.00	0.00	0.00
Q8VF84	MCG60141 OS=Mus musculus GN=Olfr1317 PE=3 SV=1	1.00	1.00	0.08	1.00
Q8VfZ5	MCG141584 OS=Mus musculus GN=Olfr516 PE=3 SV=1	1.00	0.04	1.00	1.00
Q8VHM5	Heterogeneous nuclear ribonucleoprotein R OS=Mus musculus GN=Hnrpr PE=2 SV=1	1.00	0.00	0.05	0.00
Q8VIJ6	Splicing factor, proline- and glutamine-rich OS=Mus musculus GN=Sfpq PE=1 SV=1	0.02	0.00	0.00	0.02
Q91V14	Solute carrier family 12 member 5 OS=Mus musculus GN=Slc12a5 PE=1 SV=2	1.00	0.01	1.00	1.00
Q91V41	Ras-related protein Rab-14 OS=Mus musculus GN=Rab14 PE=1 SV=3	1.00	0.02	1.00	1.00
Q91V55	Ribosomal protein S5 OS=Mus musculus GN=Rps5 PE=2 SV=1	1.00	1.00	1.00	0.02
Q91V64	Isochorismatase domain-containing protein 1 OS=Mus musculus GN=Isoc1 PE=2 SV=1	1.00	1.00	1.00	0.02
Q91VB8	Alpha globin 1 OS=Mus musculus GN=Hba-a1 PE=2 SV=1	0.02	0.02	0.00	0.00
Q91V17	Ribonuclease inhibitor OS=Mus musculus GN=Rnh1 PE=1 SV=1	1.00	1.00	0.00	0.02
Q91VM5	Heterogeneous nuclear ribonucleoprotein G-like 1 OS=Mus musculus GN=Rbmxl1 PE=2 SV=1	1.00	0.08	0.02	0.00
Q91VM9	Inorganic pyrophosphatase 2, mitochondrial OS=Mus musculus GN=Ppa2 PE=2 SV=1	1.00	1.00	0.08	1.00
Q91W90	Thioredoxin domain-containing protein 5 OS=Mus musculus GN=Txndc5 PE=1 SV=2	1.00	0.00	0.02	0.00
Q91WJ8	Far upstream element-binding protein 1 OS=Mus musculus GN=Fubp1 PE=1 SV=1	1.00	0.03	1.00	0.02
Q91WQ3	Tyrosyl-tRNA synthetase, cytoplasmic OS=Mus musculus GN=Yars PE=2 SV=3	1.00	1.00	0.05	1.00
Q91X79	Elastase 1, pancreatic OS=Mus musculus GN=Cela1 PE=2 SV=1	1.00	0.00	0.02	0.00
Q91XV3	Brain acid soluble protein 1 OS=Mus musculus GN=Basp1 PE=1 SV=3	0.00	0.00	0.00	0.00
Q91Y47	Coagulation factor XI OS=Mus musculus GN=F11 PE=2 SV=2	1.00	0.03	1.00	1.00

Q91YD9	Neural Wiskott-Aldrich syndrome protein OS=Mus musculus GN=Wasl PE=1 SV=1	1.00	1.00	0.05	1.00
Q91YQ5	Dolichyl-diphosphooligosaccharide--protein glycosyltransferase subunit 1 OS=Mus musculus GN=Rpn1 PE=2 SV=1	1.00	0.04	0.03	0.00
Q921B4	Neurotrophin receptor-interacting factor 2 OS=Mus musculus GN=Nrf2 PE=1 SV=1	0.03	0.04	0.05	0.03
Q921F2	TAR DNA-binding protein 43 OS=Mus musculus GN=Tardbp PE=1 SV=1	1.00	0.02	1.00	0.02
Q92111	Serotransferrin OS=Mus musculus GN=Tf PE=1 SV=1	1.00	0.07	1.00	1.00
Q922F4	Tubulin beta-6 chain OS=Mus musculus GN=Tubb6 PE=1 SV=1	1.00	1.00	1.00	0.04
Q922I7	MCG13402, isoform CRA_c OS=Mus musculus GN=Ptpb1 PE=2 SV=1	0.02	0.00	0.00	0.02
Q922U2	Keratin, type II cytoskeletal 5 OS=Mus musculus GN=Krt5 PE=1 SV=1	0.02	0.00	0.00	0.00
Q923D4	Splicing factor 3B subunit 5 OS=Mus musculus GN=Sf3b5 PE=2 SV=1	1.00	0.02	1.00	1.00
Q93092	Transaldolase OS=Mus musculus GN=Taldo1 PE=1 SV=2	1.00	0.04	1.00	0.00
Q99020	Heterogeneous nuclear ribonucleoprotein A/B OS=Mus musculus GN=Hnmpab PE=1 SV=1	1.00	0.00	1.00	0.00
Q99J77	N-acetylneuraminic acid synthase (Sialic acid synthase) OS=Mus musculus GN=Nans PE=2 SV=1	1.00	1.00	1.00	0.02
Q99JF8	PC4 and SFRS1-interacting protein OS=Mus musculus GN=Pspj1 PE=1 SV=1	1.00	1.00	0.04	1.00
Q99JI6	Ras-related protein Rap-1b OS=Mus musculus GN=Rap1b PE=2 SV=2	1.00	0.02	0.02	0.02
Q99JP7	Gamma-glutamyltransferase 7 OS=Mus musculus GN=Ggt7 PE=1 SV=2	1.00	0.07	1.00	1.00
Q99JY9	Actin-related protein 3 OS=Mus musculus GN=Actr3 PE=1 SV=3	1.00	1.00	1.00	0.00
Q99K48	Non-POU domain-containing octamer-binding protein OS=Mus musculus GN=Nono PE=1 SV=3	1.00	0.00	0.03	1.00
Q99KC8	von Willebrand factor A domain-containing protein 5A OS=Mus musculus GN=Vwa5a PE=1 SV=2	1.00	1.00	1.00	0.02
Q99KF0	Caspase recruitment domain-containing protein 14 OS=Mus musculus GN=Card14 PE=2 SV=2	1.00	1.00	0.03	1.00
Q99KF1	Transmembrane emp24 domain-containing protein 9 OS=Mus musculus GN=Tmed9 PE=2 SV=2	1.00	0.00	0.02	0.00
Q99K10	Aconitate hydratase, mitochondrial OS=Mus musculus GN=Aco2 PE=1 SV=1	1.00	1.00	0.02	0.04
Q99KJ8	Dynactin subunit 2 OS=Mus musculus GN=Dctn2 PE=1 SV=3	1.00	1.00	1.00	0.02
Q99KV1	DnaJ homolog subfamily B member 11 OS=Mus musculus GN=Dnajb11 PE=1 SV=1	1.00	1.00	1.00	0.04
Q99L47	Hsc70-interacting protein OS=Mus musculus GN=St13 PE=2 SV=1	1.00	1.00	1.00	0.02
Q99LC5	Electron transfer flavoprotein subunit alpha, mitochondrial OS=Mus musculus GN=Etfa PE=1 SV=2	1.00	0.02	1.00	0.02
Q99LD8	N(G),N(G)-dimethylarginine dimethylaminohydrolase 2 OS=Mus musculus GN=Ddah2 PE=1 SV=1	1.00	1.00	0.02	1.00
Q99LJ6	Glutathione peroxidase 7 OS=Mus musculus GN=Gpx7 PE=2 SV=1	1.00	0.08	1.00	1.00
Q99LX0	Protein DJ-1 OS=Mus musculus GN=Park7 PE=1 SV=1	1.00	1.00	0.04	0.00
Q99M74	Keratin, type II cuticular Hb2 OS=Mus musculus GN=Krt82 PE=2 SV=2	1.00	0.03	0.06	1.00
Q99MD9	Nuclear autoantigenic sperm protein OS=Mus musculus GN=Nasp PE=1 SV=2	1.00	0.00	0.03	1.00
Q99MR6	Serrate RNA effector molecule homolog OS=Mus musculus GN=Srrt PE=1 SV=1	0.06	1.00	1.00	1.00
Q99P72	Reticulon-4 OS=Mus musculus GN=Rtn4 PE=1 SV=2	1.00	0.02	1.00	0.00
Q99PL5	Ribosome-binding protein 1 OS=Mus musculus GN=Rrbp1 PE=2 SV=2	1.00	0.00	0.00	0.00
Q99PS0	Keratin, type I cytoskeletal 23 OS=Mus musculus GN=Krt23 PE=2 SV=1	1.00	1.00	1.00	0.02
Q99PT1	Rho GDP-dissociation inhibitor 1 OS=Mus musculus GN=Arhgdia PE=1 SV=3	0.03	0.00	1.00	0.00
Q9CPN9	RIKEN cDNA 2210010C04, isoform CRA_b OS=Mus musculus GN=2210010C04Rik PE=2 SV=1	1.00	0.06	0.03	0.00
Q9CPQ1	Cytochrome c oxidase subunit 6C OS=Mus musculus GN=Cox6c PE=1 SV=3	0.04	0.02	1.00	1.00
Q9CPQ5	Centromere protein Q OS=Mus musculus GN=Cenpq PE=2 SV=1	1.00	1.00	1.00	0.04
Q9CPR4	60S ribosomal protein L17 OS=Mus musculus GN=Rpl17 PE=2 SV=3	1.00	0.02	0.00	0.00
Q9CPV4	Glyoxalase domain-containing protein 4 OS=Mus musculus GN=Glod4 PE=2 SV=1	1.00	1.00	0.08	1.00
Q9CPW4	Actin-related protein 2/3 complex subunit 5 OS=Mus musculus GN=Arpc5 PE=2 SV=3	1.00	1.00	0.02	1.00
Q9CQ52	Elastase 3, pancreatic OS=Mus musculus GN=Cela3b PE=2 SV=1	1.00	0.00	0.00	0.00
Q9CQ65	S-methyl-5'-thioadenosine phosphorylase OS=Mus musculus GN=Mtap PE=2 SV=1	1.00	1.00	0.02	1.00
Q9CQC2	Colipase OS=Mus musculus GN=Clips PE=2 SV=1	1.00	0.00	1.00	1.00
Q9CQF9	Prenylcysteine oxidase OS=Mus musculus GN=Pcyox1 PE=1 SV=1	1.00	1.00	1.00	0.02
Q9CQI6	Coactosin-like protein OS=Mus musculus GN=Cott1 PE=1 SV=3	1.00	0.07	1.00	1.00
Q9CQI7	ATP synthase subunit b, mitochondrial OS=Mus musculus GN=Atp5f1 PE=1 SV=1	1.00	1.00	0.05	1.00
Q9CQR2	40S ribosomal protein S21 OS=Mus musculus GN=Rps21 PE=2 SV=1	1.00	0.00	0.00	0.00
Q9CQV8	14-3-3 protein beta/alpha OS=Mus musculus GN=Ywhab PE=1 SV=3	0.02	0.00	0.00	0.00
Q9CR35	Chymotrypsinogen B OS=Mus musculus GN=Ctrb1 PE=2 SV=1	0.00	0.00	0.00	0.00
Q9CR57	60S ribosomal protein L14 OS=Mus musculus GN=Rpl14 PE=2 SV=3	1.00	0.00	0.00	0.00
Q9CRB6	Tubulin polymerization-promoting protein family member 3 OS=Mus musculus GN=Tppp3 PE=1 SV=1	1.00	0.02	0.00	0.00
Q9CS72	Filamin-A-interacting protein 1 OS=Mus musculus GN=Filip1 PE=1 SV=2	0.07	0.08	1.00	1.00
Q9CWF2	Tubulin beta-2B chain OS=Mus musculus GN=Tubb2b PE=1 SV=1	1.00	0.00	0.02	0.00
Q9CWZ3	RNA-binding protein 8A OS=Mus musculus GN=Rbm8a PE=1 SV=3	1.00	0.02	1.00	1.00
Q9CX80	Cytoglobin OS=Mus musculus GN=Cygb PE=2 SV=1	1.00	1.00	1.00	0.02
Q9CX86	Uncharacterized protein OS=Mus musculus GN=Hnmpa0 PE=2 SV=1	1.00	1.00	1.00	0.02
Q9CXW4	60S ribosomal protein L11 OS=Mus musculus GN=Rpl11 PE=1 SV=4	1.00	0.04	1.00	0.02
Q9CY50	Translocon-associated protein subunit alpha OS=Mus musculus GN=Ssr1 PE=1 SV=1	0.06	0.00	0.02	0.00
Q9CY58	Plasminogen activator inhibitor 1 RNA-binding protein OS=Mus musculus GN=Serbp1 PE=1 SV=2	0.07	0.00	1.00	0.00
Q9CYZ2	Tumor protein D54 OS=Mus musculus GN=Tpd52l2 PE=1 SV=1	1.00	1.00	0.02	0.04
Q9CZ13	Cytochrome b-c1 complex subunit 1, mitochondrial OS=Mus musculus GN=Uqcrc1 PE=1 SV=2	1.00	1.00	0.02	1.00
Q9CZ69	CKLF-like MARVEL transmembrane domain-containing protein 6 OS=Mus musculus GN=Cmtm6 PE=2 SV=1	1.00	1.00	1.00	0.04
Q9CZM2	60S ribosomal protein L15 OS=Mus musculus GN=Rpl15 PE=2 SV=4	1.00	1.00	0.07	1.00
Q9CZN7	Serine hydroxymethyltransferase OS=Mus musculus GN=Shtmt2 PE=2 SV=1	1.00	0.06	1.00	1.00
Q9CZX8	40S ribosomal protein S19 OS=Mus musculus GN=Rps19 PE=1 SV=3	0.00	0.00	0.02	0.00
Q9D051	Pyruvate dehydrogenase E1 component subunit beta, mitochondrial OS=Mus musculus GN=Pdhb PE=1 SV=1	1.00	1.00	0.06	1.00
Q9D0E1	Heterogeneous nuclear ribonucleoprotein M OS=Mus musculus GN=Hnmpm PE=1 SV=3	1.00	0.08	0.02	0.07
Q9D0F9	Phosphoglucomutase-1 OS=Mus musculus GN=Pgm1 PE=1 SV=4	1.00	0.08	0.02	1.00
Q9D0J8	Parathyrosin OS=Mus musculus GN=Ptms PE=2 SV=3	0.02	0.02	0.02	0.02
Q9D0K2	Succinyl-CoA:3-ketoacid-coenzyme A transferase 1, mitochondrial OS=Mus musculus GN=Oxct1 PE=1 SV=1	1.00	0.02	1.00	1.00

Q9D0T1	NHP2-like protein 1 OS=Mus musculus GN=Nhp2l1 PE=2 SV=4	1.00	1.00	0.04	1.00
Q9D1D4	Transmembrane emp24 domain-containing protein 10 OS=Mus musculus GN=Tmed10 PE=2 SV=1	1.00	0.00	1.00	0.00
Q9D1D6	Collagen triple helix repeat-containing protein 1 OS=Mus musculus GN=Cthrc1 PE=2 SV=1	1.00	0.05	1.00	1.00
Q9D1J3	SAP domain-containing ribonucleoprotein OS=Mus musculus GN=Sarnp PE=1 SV=3	1.00	0.03	0.02	1.00
Q9D1M0	Protein SEC13 homolog OS=Mus musculus GN=Sec13 PE=2 SV=3	1.00	1.00	0.02	1.00
Q9D1R9	60S ribosomal protein L34 OS=Mus musculus GN=Rpl34 PE=3 SV=2	1.00	0.02	0.02	0.06
Q9D281	Protein Noxp20 OS=Mus musculus GN=Fam114a1 PE=2 SV=1	1.00	1.00	1.00	0.02
Q9D2M8	Ubiquitin-conjugating enzyme E2 variant 2 OS=Mus musculus GN=Ube2v2 PE=2 SV=4	1.00	0.02	0.02	0.04
Q9D358	Low molecular weight phosphotyrosine protein phosphatase OS=Mus musculus GN=Acp1 PE=2 SV=3	1.00	0.04	1.00	1.00
Q9D3D9	ATP synthase subunit delta, mitochondrial OS=Mus musculus GN=Atp5d PE=1 SV=1	0.02	0.00	0.03	1.00
Q9D662	Protein transport protein Sec23B OS=Mus musculus GN=Sec23b PE=2 SV=1	1.00	1.00	1.00	0.02
Q9D6N5	Dr1-associated corepressor OS=Mus musculus GN=Drap1 PE=2 SV=3	1.00	0.05	1.00	1.00
Q9D773	39S ribosomal protein L2, mitochondrial OS=Mus musculus GN=Mrlp2 PE=2 SV=1	1.00	1.00	0.03	0.08
Q9D8B3	Charged multivesicular body protein 4b OS=Mus musculus GN=Chmp4b PE=2 SV=2	1.00	0.02	0.02	0.02
Q9D8E6	60S ribosomal protein L4 OS=Mus musculus GN=Rpl4 PE=1 SV=3	1.00	1.00	0.02	0.02
Q9D8N0	Elongation factor 1-gamma OS=Mus musculus GN=Eef1g PE=1 SV=3	1.00	0.02	0.00	0.04
Q9D8U4	Complement C1q tumor necrosis factor-related protein 2 OS=Mus musculus GN=C1qtnf2 PE=2 SV=1	1.00	0.02	0.03	1.00
Q9DAK9	14 kDa phosphohistidine phosphatase OS=Mus musculus GN=Phpt1 PE=1 SV=1	1.00	0.04	1.00	1.00
Q9DAW9	Calponin-3 OS=Mus musculus GN=Cnn3 PE=2 SV=1	1.00	0.00	1.00	1.00
Q9DB77	Cytochrome b-c1 complex subunit 2, mitochondrial OS=Mus musculus GN=Uqcrc2 PE=1 SV=1	0.04	0.02	0.02	1.00
Q9DBG5	Perilipin-3 OS=Mus musculus GN=Plin3 PE=1 SV=1	1.00	1.00	0.03	1.00
Q9DBG6	Dolichyl-diphosphooligosaccharide--protein glycosyltransferase subunit 2 OS=Mus musculus GN=Rpn2 PE=2 SV=1	0.03	1.00	1.00	0.02
Q9DBJ1	Phosphoglycerate mutase 1 OS=Mus musculus GN=Pgam1 PE=1 SV=3	0.00	0.00	0.00	0.00
Q9DBP5	UMP-CMP kinase OS=Mus musculus GN=Cmpk1 PE=1 SV=1	1.00	1.00	0.05	1.00
Q9DBZ1	Inhibitor of nuclear factor kappa-B kinase-interacting protein OS=Mus musculus GN=Ikkip PE=1 SV=2	1.00	1.00	1.00	0.07
Q9DC29	ATP-binding cassette sub-family B member 6, mitochondrial OS=Mus musculus GN=Abcb6 PE=1 SV=1	1.00	0.03	1.00	1.00
Q9DCD0	6-phosphogluconate dehydrogenase, decarboxylating OS=Mus musculus GN=Pgd PE=2 SV=3	1.00	1.00	0.02	0.02
Q9DCF9	Translocon-associated protein subunit gamma OS=Mus musculus GN=Ssr3 PE=1 SV=1	1.00	1.00	1.00	0.02
Q9DCL9	Multifunctional protein ADE2 OS=Mus musculus GN=Paics PE=1 SV=4	1.00	1.00	1.00	0.02
Q9DCN2	NADH-cytochrome b5 reductase 3 OS=Mus musculus GN=Cyb5r3 PE=1 SV=3	1.00	0.04	0.00	0.00
Q9DCQ2	Putative L-aspartate dehydrogenase OS=Mus musculus GN=Aspdh PE=1 SV=1	1.00	0.04	0.02	1.00
Q9DCV7	Keratin, type II cytoskeletal 7 OS=Mus musculus GN=Krt7 PE=1 SV=1	1.00	1.00	1.00	0.00
Q9DCW4	Electron transfer flavoprotein subunit beta OS=Mus musculus GN=Etfb PE=1 SV=3	1.00	0.00	0.00	0.00
Q9DCX2	ATP synthase subunit d, mitochondrial OS=Mus musculus GN=Atp5h PE=1 SV=3	1.00	1.00	0.02	1.00
Q9EPC2	Fibroblast growth factor 23 OS=Mus musculus GN=Fgf23 PE=1 SV=1	1.00	0.03	1.00	1.00
Q9EQU5	Protein SET OS=Mus musculus GN=Set PE=1 SV=1	0.02	0.00	0.02	0.00
Q9ER05	Chymopasin OS=Mus musculus GN=Ctr1 PE=2 SV=1	1.00	0.02	0.02	0.00
Q9EST1	Gasdermin-A OS=Mus musculus GN=Gsdma PE=2 SV=1	1.00	1.00	0.06	0.00
Q9EST5	Acidic leucine-rich nuclear phosphoprotein 32 family member B OS=Mus musculus GN=Anp32b PE=1 SV=1	1.00	0.02	0.03	1.00
Q9JHJ0	Tropomodulin-3 OS=Mus musculus GN=Tmod3 PE=1 SV=1	1.00	1.00	1.00	0.05
Q9JHU4	Cytoplasmic dynein 1 heavy chain 1 OS=Mus musculus GN=Dync1h1 PE=1 SV=2	1.00	1.00	1.00	0.02
Q9JHU9	Inositol-3-phosphate synthase 1 OS=Mus musculus GN=Isyna1 PE=2 SV=1	0.07	1.00	1.00	1.00
Q9JI16	Alcohol dehydrogenase [NADP+] OS=Mus musculus GN=Akr1a1 PE=1 SV=3	1.00	0.00	0.04	0.07
Q9JIW9	Ras-related protein Ral-B OS=Mus musculus GN=Ralb PE=2 SV=1	1.00	0.05	1.00	1.00
Q9JJG7	Calcium-binding protein 8 OS=Mus musculus GN=Caln1 PE=2 SV=1	1.00	0.02	1.00	1.00
Q9JJU8	SH3 domain-binding glutamic acid-rich-like protein OS=Mus musculus GN=Sh3bgl1 PE=2 SV=1	1.00	0.02	0.02	0.02
Q9JJW6	RNA and export factor-binding protein 2 OS=Mus musculus GN=Refbp2 PE=1 SV=1	1.00	0.02	0.02	0.06
Q9JK81	UPF0160 protein MYG1, mitochondrial OS=Mus musculus GN=Myg1 PE=2 SV=1	1.00	0.03	1.00	1.00
Q9JK88	Serpin I2 OS=Mus musculus GN=Serpini2 PE=1 SV=1	1.00	0.03	1.00	1.00
Q9JLJ2	4-trimethylaminobutyraldehyde dehydrogenase OS=Mus musculus GN=Aldh9a1 PE=1 SV=1	1.00	0.03	1.00	1.00
Q9JMA4	Killer cell lectin-like receptor, subfamily A, member 17 OS=Mus musculus GN=Klra17 PE=2 SV=1	1.00	1.00	0.02	1.00
Q9JMG7	Hepatoma-derived growth factor-related protein 3 OS=Mus musculus GN=Hdgrp3 PE=1 SV=2	1.00	1.00	0.06	1.00
Q9QU10	Transforming protein RhoA OS=Mus musculus GN=Rhoa PE=1 SV=1	1.00	0.04	0.00	0.02
Q9QUM9	Proteasome subunit alpha type-6 OS=Mus musculus GN=Psm6 PE=1 SV=1	1.00	1.00	0.02	0.02
Q9QXL1	Kinesin-like protein KIF21B OS=Mus musculus GN=Kif21b PE=1 SV=2	1.00	1.00	0.08	1.00
Q9QXS1	Plectin OS=Mus musculus GN=Plec PE=1 SV=2	1.00	1.00	0.04	1.00
Q9QXT0	Protein canopy homolog 2 OS=Mus musculus GN=Cnpy2 PE=2 SV=1	1.00	0.03	1.00	0.02
Q9QYB1	Chloride intracellular channel protein 4 OS=Mus musculus GN=Clic4 PE=1 SV=3	1.00	1.00	1.00	0.02
Q9QYB5	Gamma-adducin OS=Mus musculus GN=Add3 PE=1 SV=2	1.00	1.00	1.00	0.00
Q9QYC0	Alpha-adducin OS=Mus musculus GN=Add1 PE=1 SV=2	1.00	1.00	0.00	0.00
Q9QZD9	Eukaryotic translation initiation factor 3 subunit I OS=Mus musculus GN=Eif3i PE=1 SV=1	1.00	0.08	1.00	1.00
Q9R0N0	Galactokinase OS=Mus musculus GN=Galk1 PE=2 SV=1	1.00	0.03	1.00	1.00
Q9R0P3	S-formylglutathione hydrolase OS=Mus musculus GN=Esdl PE=2 SV=1	1.00	1.00	0.05	1.00
Q9R0P5	Dextrin OS=Mus musculus GN=Dstn PE=1 SV=3	1.00	1.00	1.00	0.00
Q9R0P6	Signal peptidase complex catalytic subunit SEC11A OS=Mus musculus GN=Sec11a PE=2 SV=1	1.00	0.03	1.00	1.00
Q9R0Q3	Transmembrane emp24 domain-containing protein 2 OS=Mus musculus GN=Tmed2 PE=1 SV=1	0.02	0.00	0.02	0.03
Q9R0T7	MCG15085 OS=Mus musculus GN=Try4 PE=2 SV=1	1.00	1.00	1.00	0.00
Q9R0T8	Inhibitor of nuclear factor kappa-B kinase subunit epsilon OS=Mus musculus GN=Ikbe PE=2 SV=2	1.00	0.03	0.04	0.05
Q9R0Y5	Adenylate kinase isoenzyme 1 OS=Mus musculus GN=Ak1 PE=1 SV=1	1.00	1.00	1.00	0.03
Q9R1P0	Proteasome subunit alpha type-4 OS=Mus musculus GN=Psm4 PE=1 SV=1	1.00	0.00	1.00	1.00
Q9R1P3	Proteasome subunit beta type-2 OS=Mus musculus GN=Psm2 PE=1 SV=1	1.00	0.02	0.02	0.02

Q9R1P4	Proteasome subunit alpha type-1 OS=Mus musculus GN=Psm1 PE=1 SV=1	1.00	0.00	0.07	1.00
Q9WTL8	Aryl hydrocarbon receptor nuclear translocator-like protein 1 OS=Mus musculus GN=Arntl PE=1 SV=2	1.00	1.00	1.00	0.07
Q9WTP6	Adenylate kinase 2, mitochondrial OS=Mus musculus GN=Ak2 PE=1 SV=5	1.00	0.00	1.00	1.00
Q9WTQ5	A-kinase anchor protein 12 OS=Mus musculus GN=Akap12 PE=1 SV=1	1.00	0.00	0.00	0.00
Q9WTX5	S-phase kinase-associated protein 1 OS=Mus musculus GN=Skp1 PE=1 SV=3	1.00	0.02	0.02	1.00
Q9WUK2	Eukaryotic translation initiation factor 4H OS=Mus musculus GN=Eif4h PE=1 SV=3	1.00	1.00	1.00	0.02
Q9WUM4	Coronin-1C OS=Mus musculus GN=Coro1c PE=1 SV=2	1.00	1.00	1.00	0.02
Q9WUU7	Cathepsin Z OS=Mus musculus GN=Ctsz PE=2 SV=1	1.00	0.02	1.00	1.00
Q9WV80	Sorting nexin-1 OS=Mus musculus GN=Snx1 PE=1 SV=1	1.00	1.00	0.02	0.02
Q9WVA4	Transgelin-2 OS=Mus musculus GN=Tagln2 PE=1 SV=4	1.00	0.00	0.00	0.00
Q9WVK4	EH domain-containing protein 1 OS=Mus musculus GN=Ehd1 PE=1 SV=1	1.00	1.00	1.00	0.00
Q9Z0E6	Interferon-induced guanylate-binding protein 2 OS=Mus musculus GN=Gbp2 PE=1 SV=1	1.00	1.00	1.00	0.00
Q9Z0X1	Apoptosis-inducing factor 1, mitochondrial OS=Mus musculus GN=Aifm1 PE=1 SV=1	1.00	0.00	1.00	0.06
Q9Z1N5	Spliceosome RNA helicase Ddx39b OS=Mus musculus GN=Ddx39b PE=1 SV=1	0.05	0.03	1.00	1.00
Q9Z1Q5	Chloride intracellular channel protein 1 OS=Mus musculus GN=Clic1 PE=1 SV=3	1.00	0.00	0.02	0.02
Q9Z1Q9	Valyl-tRNA synthetase OS=Mus musculus GN=Vars PE=2 SV=1	1.00	0.02	0.02	1.00
Q9Z1R9	MCG124046 OS=Mus musculus GN=Prss1 PE=2 SV=1	1.00	0.02	1.00	0.02
Q9Z1T2	Thrombospondin-4 OS=Mus musculus GN=Thbs4 PE=2 SV=1	1.00	0.05	1.00	1.00
Q9Z1Z2	Serine-threonine kinase receptor-associated protein OS=Mus musculus GN=Strap PE=1 SV=2	1.00	0.02	1.00	1.00
Q9Z204	Heterogeneous nuclear ribonucleoproteins C1/C2 OS=Mus musculus GN=Hnrnpc PE=1 SV=1	1.00	0.00	1.00	0.00
Q9Z2M7	Phosphomannomutase 2 OS=Mus musculus GN=Pmm2 PE=2 SV=1	1.00	1.00	0.06	1.00
Q9Z2U0	Proteasome subunit alpha type-7 OS=Mus musculus GN=Psm7 PE=1 SV=1	1.00	0.00	0.00	0.00
Q9Z2U1	Proteasome subunit alpha type-5 OS=Mus musculus GN=Psm5 PE=1 SV=1	1.00	1.00	1.00	0.02
Q9Z2W0	Aspartyl aminopeptidase OS=Mus musculus GN=Dnpep PE=2 SV=2	1.00	1.00	0.02	1.00
Q9Z2X1	Heterogeneous nuclear ribonucleoprotein F OS=Mus musculus GN=Hnrnpf PE=1 SV=3	1.00	0.02	1.00	0.02

Unique Peptides				Spectral Counts				Normalized Spectral Counts				gene
e15.5	e17.5	p2	p14	e15.5	e17.5	p2	p14	e15.5	e17.5	p2	p14	gene
0	0	1	0	0	0	0	1	0	0.0000	0.0000	0.0003	0.0000 Tbc1d16
0	1	0	0	0	0	2	0	0	0.0000	0.0003	0.0000	0.0000 Fsip2
1	0	0	0	0	1	0	0	0	0.0008	0.0000	0.0000	0.0000 Agrn
0	5	3	10	0	23	14	47	0.0000	0.0032	0.0040	0.0132 Hbb-b1	
0	11	7	8	0	92	32	18	0.0000	0.0129	0.0091	0.0051 Cpb1	
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0003 Zcchc11	
0	1	1	1	0	2	4	2	0.0000	0.0003	0.0011	0.0006 Tpm1	
1	1	1	0	1	6	2	0	0.0008	0.0008	0.0006	0.0000 Gm9858	
1	0	0	0	3	0	0	0	0.0023	0.0000	0.0000	0.0000 Stard13	
0	1	3	3	0	1	6	4	0.0000	0.0001	0.0017	0.0011 Pdia2	
0	1	0	0	0	1	0	0	0.0000	0.0001	0.0000	0.0000 4930485B16Rik	
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0003 Gstm5	
0	0	0	1	0	0	0	2	0.0000	0.0000	0.0000	0.0006 Ccdc144b	
0	1	0	1	0	1	0	2	0.0000	0.0001	0.0000	0.0006 Ryr1	
0	0	1	0	0	0	1	0	0.0000	0.0000	0.0003	0.0000 Olfr288	
0	3	0	0	0	9	0	0	0.0000	0.0013	0.0000	0.0000 5430421N21Rik	
0	3	8	35	0	12	22	74	0.0000	0.0017	0.0062	0.0208 Spna2	
0	1	1	3	0	2	3	3	0.0000	0.0003	0.0009	0.0008 Tpm1	
0	2	3	19	0	3	5	48	0.0000	0.0004	0.0014	0.0135 Dsp	
0	14	11	44	0	46	33	99	0.0000	0.0065	0.0094	0.0279 Ahnak	
1	1	1	0	1	1	1	0	0.0008	0.0001	0.0003	0.0000 Urb2	
0	1	0	0	0	6	0	0	0.0000	0.0008	0.0000	0.0000 Fcgbp	
1	1	0	0	11	3	0	0	0.0085	0.0004	0.0000	0.0000 Upf2	
0	1	0	0	0	2	0	0	0.0000	0.0003	0.0000	0.0000 D1Ert622e	
1	1	1	0	5	6	2	0	0.0039	0.0008	0.0006	0.0000 ncharacterized	
0	0	1	0	0	0	1	0	0.0000	0.0000	0.0003	0.0000 Hpn	
0	0	0	4	0	0	0	6	0.0000	0.0000	0.0000	0.0017 Ahnak2	
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0003 Sec22b	
0	1	1	7	0	14	6	12	0.0000	0.0020	0.0017	0.0034 Dpysl2	
0	0	1	0	0	0	1	0	0.0000	0.0000	0.0003	0.0000 Dld	
1	2	0	1	2	2	0	2	0.0015	0.0003	0.0000	0.0006 Prkcxh	
0	2	1	0	0	5	6	0	0.0000	0.0007	0.0017	0.0000 Prdx4	
0	0	0	2	0	0	0	7	0.0000	0.0000	0.0000	0.0020 Ptgis	
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0003 Dpysl4	
0	1	1	0	0	7	2	0	0.0000	0.0010	0.0006	0.0000 Phb2	
0	0	1	0	0	0	2	0	0.0000	0.0000	0.0006	0.0000 Srsf5	
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0003 Capn1	
0	3	0	0	0	5	0	0	0.0000	0.0007	0.0000	0.0000 Anp32a	
0	0	0	4	0	0	0	7	0.0000	0.0000	0.0000	0.0020 Anxa3	
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0003 Anxa8	
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0003 Myadm	
0	1	0	0	0	2	0	0	0.0000	0.0003	0.0000	0.0000 Nudc	
0	1	0	0	0	2	0	0	0.0000	0.0003	0.0000	0.0000 Bcat2	
0	3	3	3	0	7	8	6	0.0000	0.0010	0.0023	0.0017 Calu	
0	0	1	8	0	0	6	25	0.0000	0.0000	0.0017	0.0070 Pitr	
0	1	0	2	0	2	0	4	0.0000	0.0003	0.0000	0.0011 Ddost	
1	0	0	0	2	0	0	0	0.0015	0.0000	0.0000	0.0000 Aox1	
0	0	1	0	0	0	2	0	0.0000	0.0000	0.0006	0.0000 Hmgb3	
1	0	0	0	1	0	0	0	0.0008	0.0000	0.0000	0.0000 Banf1	
0	1	0	0	0	3	0	0	0.0000	0.0004	0.0000	0.0000 Pgrmc1	
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0003 Impa1	
0	0	0	2	0	0	0	2	0.0000	0.0000	0.0000	0.0006 Entpd2	
0	0	0	1	0	0	0	2	0.0000	0.0000	0.0000	0.0006 Copb2	
0	0	3	1	0	0	6	1	0.0000	0.0000	0.0017	0.0003 7-Sep	
0	0	1	1	0	0	4	2	0.0000	0.0000	0.0011	0.0006 Rpl35a	
0	1	2	0	0	4	5	0	0.0000	0.0006	0.0014	0.0000 Psmb5	
1	2	0	2	2	12	0	3	0.0015	0.0017	0.0000	0.0008 Eef1b	
0	1	0	0	0	1	0	0	0.0000	0.0001	0.0000	0.0000 Epb41I2	
0	1	1	0	0	8	2	0	0.0000	0.0011	0.0006	0.0000 Psma3	
0	0	1	0	0	0	1	0	0.0000	0.0000	0.0003	0.0000 Sfn	
0	2	2	2	0	9	8	4	0.0000	0.0013	0.0023	0.0011 Ugdh	
0	0	1	0	0	0	2	0	0.0000	0.0000	0.0006	0.0000 Snx3	
0	0	1	0	0	0	2	0	0.0000	0.0000	0.0006	0.0000 Wdr1	
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0003 Capns1	
0	2	0	0	0	5	0	0	0.0000	0.0007	0.0000	0.0000 Dync1i2	
7	7	6	7	37	88	28	19	0.0285	0.0124	0.0079	0.0054 Hnrmpa2b1	
0	2	2	3	0	4	5	5	0.0000	0.0006	0.0014	0.0014 Idh1	
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0003 Cyp8b1	
1	1	1	0	1	11	4	0	0.0008	0.0015	0.0011	0.0000 Rbm3	

0	1	1	0	0	1	4	0	0.0000	0.0001	0.0011	0.0000	Rnase1
0	22	8	11	0	324	68	41	0.0000	0.0456	0.0193	0.0115	Amy2
0	1	0	1	0	1	0	10	0.0000	0.0001	0.0000	0.0028	Krt1
0	0	0	3	0	0	0	6	0.0000	0.0000	0.0000	0.0017	Fabp4
1	9	4	2	2	35	15	4	0.0015	0.0049	0.0043	0.0011	Aldoa
0	0	1	0	0	0	1	0	0.0000	0.0000	0.0003	0.0000	Got2
0	5	2	6	0	58	13	15	0.0000	0.0082	0.0037	0.0042	Cela2a
2	0	0	6	6	0	0	10	0.0046	0.0000	0.0000	0.0028	Tuba1b
1	3	2	3	3	25	5	4	0.0023	0.0035	0.0014	0.0011	Ldha
0	0	2	1	0	0	9	1	0.0000	0.0000	0.0026	0.0003	Gpi
0	2	1	3	0	14	11	5	0.0000	0.0020	0.0031	0.0014	Prss2
2	6	4	14	11	60	40	66	0.0085	0.0084	0.0114	0.0186	Anxa2
0	1	0	4	0	8	0	6	0.0000	0.0011	0.0000	0.0017	Alb
3	4	3	3	10	30	16	8	0.0077	0.0042	0.0045	0.0023	Hsp90aa1
3	5	4	4	6	24	9	6	0.0046	0.0034	0.0026	0.0017	Pdia4
4	10	6	15	10	58	23	43	0.0077	0.0082	0.0065	0.0121	Hsp90b1
0	1	1	2	0	7	7	6	0.0000	0.0010	0.0020	0.0017	S100a10
1	4	3	4	6	34	20	15	0.0046	0.0048	0.0057	0.0042	Sod1
0	9	3	6	0	15	23	11	0.0000	0.0021	0.0065	0.0031	Mdh2
1	0	0	1	3	0	0	1	0.0023	0.0000	0.0000	0.0003	Krt13
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Gnai2
0	1	1	1	0	2	1	1	0.0000	0.0003	0.0003	0.0003	Itgb1
8	12	7	14	37	108	44	41	0.0285	0.0152	0.0125	0.0115	P4hb
2	6	6	6	8	29	19	9	0.0062	0.0041	0.0054	0.0025	Ncl
0	1	0	1	0	1	0	1	0.0000	0.0001	0.0000	0.0003	Pgk1
0	2	0	0	0	4	0	0	0.0000	0.0006	0.0000	0.0000	Hmgn2
0	0	4	10	0	0	26	25	0.0000	0.0000	0.0074	0.0070	Anxa1
4	13	6	13	44	109	51	50	0.0340	0.0153	0.0145	0.0141	Eef1a1
1	3	2	4	2	14	7	10	0.0015	0.0020	0.0020	0.0028	Txn
0	2	2	4	0	7	15	10	0.0000	0.0010	0.0043	0.0028	Gstm1
1	2	2	3	5	16	9	9	0.0039	0.0023	0.0026	0.0025	H1f0
1	2	1	0	2	6	5	0	0.0015	0.0008	0.0014	0.0000	Sub1
1	2	2	2	1	13	7	4	0.0008	0.0018	0.0020	0.0011	Hsp90ab1
0	1	1	0	0	1	2	0	0.0000	0.0001	0.0006	0.0000	Itga5
0	2	1	2	0	5	1	2	0.0000	0.0007	0.0003	0.0006	Rpl7a
1	3	4	3	2	20	19	5	0.0015	0.0028	0.0054	0.0014	Gsn
1	1	1	1	6	10	6	4	0.0046	0.0014	0.0017	0.0011	S100a6
0	1	0	1	0	2	0	2	0.0000	0.0003	0.0000	0.0006	Rpl27a
0	2	1	1	0	4	1	1	0.0000	0.0006	0.0003	0.0003	Rps16
0	1	0	1	0	3	0	2	0.0000	0.0004	0.0000	0.0006	Rpl7
1	2	2	4	8	16	8	10	0.0062	0.0023	0.0023	0.0028	Mdh1
2	4	2	4	13	49	13	10	0.0100	0.0069	0.0037	0.0028	Rpsa
2	7	5	9	6	53	21	22	0.0046	0.0075	0.0060	0.0062	Calr
0	1	0	0	0	1	0	0	0.0000	0.0001	0.0000	0.0000	H2-Q9
2	5	5	8	3	21	15	16	0.0023	0.0030	0.0043	0.0045	Lmnb1
1	9	6	10	2	25	18	18	0.0015	0.0035	0.0051	0.0051	Anxa6
1	2	1	4	1	2	6	8	0.0008	0.0003	0.0017	0.0023	Rplp0
0	1	2	3	0	1	3	7	0.0000	0.0001	0.0009	0.0020	Gstm2
4	8	8	4	32	186	72	42	0.0247	0.0262	0.0204	0.0118	Hist1h1c
0	0	1	0	0	0	1	0	0.0000	0.0000	0.0003	0.0000	Klk1
1	6	4	7	7	37	23	40	0.0054	0.0052	0.0065	0.0113	Lgals1
1	4	1	5	2	27	9	10	0.0015	0.0038	0.0026	0.0028	Gapdh
0	0	1	0	0	0	1	0	0.0000	0.0000	0.0003	0.0000	Lamp2
2	7	4	10	14	36	26	22	0.0108	0.0051	0.0074	0.0062	Eno1
1	7	2	6	1	33	18	18	0.0008	0.0046	0.0051	0.0051	Ppia
3	4	5	6	5	45	25	16	0.0039	0.0063	0.0071	0.0045	Tpi1
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Lyz1
1	2	1	0	4	8	4	0	0.0031	0.0011	0.0011	0.0000	Pcna
0	0	0	1	0	0	0	2	0.0000	0.0000	0.0000	0.0006	Ctsd
0	0	1	1	0	0	2	2	0.0000	0.0000	0.0006	0.0006	Bsg
2	4	1	6	3	35	16	16	0.0023	0.0049	0.0045	0.0045	Cfl1
0	1	0	0	0	1	0	0	0.0000	0.0001	0.0000	0.0000	Krt19
0	1	0	3	0	1	0	7	0.0000	0.0001	0.0000	0.0020	Gstp1
0	2	1	2	0	4	6	5	0.0000	0.0006	0.0017	0.0014	Rpl13a
2	9	4	7	4	49	20	28	0.0031	0.0069	0.0057	0.0079	Serpinh1
3	13	6	16	6	51	11	44	0.0046	0.0072	0.0031	0.0124	Hspa5
0	0	1	0	0	0	1	0	0.0000	0.0000	0.0003	0.0000	Tmsb4x
0	0	1	0	0	0	2	0	0.0000	0.0000	0.0006	0.0000	Prdx3
1	7	9	25	1	35	45	105	0.0008	0.0049	0.0128	0.0296	Vim
0	0	0	6	0	0	0	12	0.0000	0.0000	0.0000	0.0034	Tgm2
0	3	3	1	0	5	16	2	0.0000	0.0007	0.0045	0.0006	Cbx3

0	1	1	1	0	1	3	1	0.0000	0.0001	0.0009	0.0003	Pnp
0	1	0	0	0	1	0	0	0.0000	0.0001	0.0000	0.0000	Cat
1	6	2	6	3	34	33	24	0.0023	0.0048	0.0094	0.0068	Ppib
1	0	0	0	2	0	0	0	0.0015	0.0000	0.0000	0.0000	Mcm3
0	1	2	2	0	11	6	2	0.0000	0.0015	0.0017	0.0006	Rps2
0	1	1	5	0	2	1	9	0.0000	0.0003	0.0003	0.0025	Tin1
0	1	1	2	0	12	4	2	0.0000	0.0017	0.0011	0.0006	Ezr
1	5	3	4	3	18	11	7	0.0023	0.0025	0.0031	0.0020	Msn
0	2	0	0	0	4	0	0	0.0000	0.0006	0.0000	0.0000	Rdx
3	6	4	3	13	51	11	6	0.0100	0.0072	0.0031	0.0017	Ptma
0	0	1	0	0	0	1	0	0.0000	0.0000	0.0003	0.0000	Glud1
0	1	0	0	0	1	0	0	0.0000	0.0001	0.0000	0.0000	Sars
5	6	4	3	33	68	24	9	0.0255	0.0096	0.0068	0.0025	Marcks
0	1	4	2	0	2	8	2	0.0000	0.0003	0.0023	0.0006	Map4
0	1	1	2	0	4	4	2	0.0000	0.0006	0.0011	0.0006	Rpl3
9	14	10	16	33	148	55	56	0.0255	0.0208	0.0156	0.0158	Pdia3
0	0	1	0	0	0	3	0	0.0000	0.0000	0.0009	0.0000	Apex1
1	0	0	0	1	0	0	0	0.0008	0.0000	0.0000	0.0000	Nap111
0	1	0	0	0	1	0	0	0.0000	0.0001	0.0000	0.0000	Nckap1
2	2	1	0	6	7	2	0	0.0046	0.0010	0.0006	0.0000	Marcks1f
0	0	0	1	0	0	0	2	0.0000	0.0000	0.0000	0.0006	Dpp4
1	0	1	0	1	0	2	0	0.0008	0.0000	0.0006	0.0000	Pabpc1
0	1	2	3	0	5	9	6	0.0000	0.0007	0.0026	0.0017	Oat
0	1	0	1	0	1	0	3	0.0000	0.0001	0.0000	0.0008	Ppic
0	1	1	0	0	1	1	0	0.0000	0.0001	0.0003	0.0000	Fkbp4
1	5	2	1	3	14	7	1	0.0023	0.0020	0.0020	0.0003	Hmgb2
0	1	4	2	0	18	10	8	0.0000	0.0025	0.0028	0.0023	Des
0	0	1	0	0	0	1	0	0.0000	0.0000	0.0003	0.0000	Dbi
2	1	1	1	5	5	1	1	0.0039	0.0007	0.0003	0.0003	Ranbp1
1	1	0	0	2	2	0	0	0.0015	0.0003	0.0000	0.0000	Mif
0	0	0	3	0	0	0	4	0.0000	0.0000	0.0000	0.0011	Rab5c
0	1	0	0	0	1	0	0	0.0000	0.0001	0.0000	0.0000	Pdha1
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Hal
0	2	0	8	0	8	0	14	0.0000	0.0011	0.0000	0.0039	Canx
2	5	4	8	4	36	20	25	0.0031	0.0051	0.0057	0.0070	Prdx1
0	3	2	4	0	21	16	10	0.0000	0.0030	0.0045	0.0028	Rpl12
0	3	2	3	0	8	12	11	0.0000	0.0011	0.0034	0.0031	Rpl18
0	1	4	2	0	1	10	4	0.0000	0.0001	0.0028	0.0011	Tagln
0	0	1	0	0	0	1	0	0.0000	0.0000	0.0003	0.0000	Hmgcl
0	1	1	1	0	1	4	2	0.0000	0.0001	0.0011	0.0006	Cap1
1	2	3	3	2	4	7	4	0.0015	0.0006	0.0020	0.0011	Tkt
0	1	0	4	0	2	0	7	0.0000	0.0003	0.0000	0.0020	
0	1	0	1	0	2	0	1	0.0000	0.0003	0.0000	0.0003	Cct8
0	0	2	3	0	0	3	6	0.0000	0.0000	0.0009	0.0017	Reg1
0	2	1	2	0	22	7	8	0.0000	0.0031	0.0020	0.0023	Hist1h1e
3	3	2	3	10	47	24	12	0.0077	0.0066	0.0068	0.0034	Hist1h1a
3	6	4	4	5	57	43	22	0.0039	0.0080	0.0122	0.0062	Hist1h1b
0	1	0	0	0	1	0	0	0.0000	0.0001	0.0000	0.0000	Hist1h1d
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Akr1b1
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Cfl2
0	1	0	0	0	4	0	0	0.0000	0.0006	0.0000	0.0000	Fkbp2
0	3	2	3	0	21	7	10	0.0000	0.0030	0.0020	0.0028	Aldh2
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Aldh3a2
0	1	1	1	0	1	2	2	0.0000	0.0001	0.0006	0.0006	Capza1
0	1	1	1	0	1	1	2	0.0000	0.0001	0.0003	0.0006	Capzb
0	1	0	0	0	2	0	0	0.0000	0.0003	0.0000	0.0000	Igfbp3
0	0	1	2	0	0	2	4	0.0000	0.0000	0.0006	0.0011	Rpl6
0	1	0	0	0	2	0	0	0.0000	0.0003	0.0000	0.0000	Rpl29
0	1	0	1	0	4	0	2	0.0000	0.0006	0.0000	0.0006	Rplp1
1	2	2	1	1	7	11	1	0.0008	0.0010	0.0031	0.0003	Rpl5
0	3	2	1	0	12	7	2	0.0000	0.0017	0.0020	0.0006	Rpl13
0	2	1	0	0	4	2	0	0.0000	0.0006	0.0006	0.0000	Rpl36
1	7	5	11	2	53	28	36	0.0015	0.0075	0.0079	0.0101	Anxa5
3	14	11	9	4	62	38	14	0.0031	0.0087	0.0108	0.0039	Lmna
0	1	0	0	0	3	0	0	0.0000	0.0004	0.0000	0.0000	Bra1
0	1	0	0	0	3	0	0	0.0000	0.0004	0.0000	0.0000	Cbr1
0	1	0	0	0	2	0	0	0.0000	0.0003	0.0000	0.0000	Cox7a2
1	1	0	0	9	7	0	0	0.0069	0.0010	0.0000	0.0000	Dnase1
4	6	3	3	27	32	11	5	0.0208	0.0045	0.0031	0.0014	Hnrmpa1
0	1	0	0	0	1	0	0	0.0000	0.0001	0.0000	0.0000	Psma2
0	0	1	2	0	0	2	9	0.0000	0.0000	0.0006	0.0025	Cav1

2-Sep

0	1	0	0	0	4	0	0	0.0000	0.0006	0.0000	0.0000	Cdkn1c
1	4	3	7	1	15	14	18	0.0008	0.0021	0.0040	0.0051	Ahcy
0	1	0	1	0	3	0	4	0.0000	0.0004	0.0000	0.0011	S100a11
2	3	4	0	5	5	9	0	0.0039	0.0007	0.0026	0.0000	Pa2g4
0	1	1	2	0	1	4	3	0.0000	0.0001	0.0011	0.0008	Rab7a
0	0	1	1	0	0	1	1	0.0000	0.0000	0.0003	0.0003	Acadl
0	0	0	2	0	0	0	3	0.0000	0.0000	0.0000	0.0008	Rpl9
0	0	1	0	0	0	0	3	0.0000	0.0000	0.0009	0.0000	Hsd17b4
0	3	1	1	0	14	4	1	0.0000	0.0020	0.0011	0.0003	Hdgf
3	10	3	10	5	62	14	21	0.0039	0.0087	0.0040	0.0059	Pkm2
0	1	0	0	0	1	0	0	0.0000	0.0001	0.0000	0.0000	Efna1
1	1	0	0	1	1	0	0	0.0008	0.0001	0.0000	0.0000	Hmga2
0	1	1	1	0	1	2	1	0.0000	0.0001	0.0006	0.0003	Rpl10a
0	3	4	6	0	13	15	12	0.0000	0.0018	0.0043	0.0034	Idh2
1	3	0	0	2	3	0	0	0.0015	0.0004	0.0000	0.0000	Stmn1
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Cyb5a
4	6	3	9	21	22	9	17	0.0162	0.0031	0.0026	0.0048	Atp5b
1	1	0	1	2	1	0	1	0.0015	0.0001	0.0000	0.0003	Ucp3
1	3	3	2	2	13	13	4	0.0015	0.0018	0.0037	0.0011	Fus
0	1	0	0	0	2	0	0	0.0000	0.0003	0.0000	0.0000	Erp29
0	1	1	2	0	1	7	3	0.0000	0.0001	0.0020	0.0008	Eef1d
0	1	1	5	0	2	3	6	0.0000	0.0003	0.0009	0.0017	Actn4
1	11	7	11	4	35	31	19	0.0031	0.0049	0.0088	0.0054	Eef2
1	2	2	2	2	14	12	3	0.0015	0.0020	0.0034	0.0008	Tpm1
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Tnks1bp1
0	1	0	0	0	1	0	0	0.0000	0.0001	0.0000	0.0000	Slc12a3
0	0	1	0	0	0	1	0	0.0000	0.0000	0.0000	0.0000	Ruvbl1
0	1	1	1	0	13	5	2	0.0000	0.0018	0.0014	0.0006	Pcbp1
0	2	1	0	0	3	3	0	0.0000	0.0004	0.0009	0.0000	Cdc42
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Cirbp
1	4	1	2	2	9	3	5	0.0015	0.0013	0.0009	0.0014	Eif4a1
1	2	2	2	1	17	11	10	0.0008	0.0024	0.0031	0.0028	Rps20
1	2	1	2	4	18	5	2	0.0031	0.0025	0.0014	0.0006	Ube2n
1	1	0	0	3	5	0	0	0.0023	0.0007	0.0000	0.0000	Drd2
2	2	1	0	5	11	1	0	0.0039	0.0015	0.0003	0.0000	Rpl26
0	1	1	1	0	2	4	2	0.0000	0.0003	0.0011	0.0006	Rpl27
1	1	1	1	1	13	4	2	0.0008	0.0018	0.0011	0.0006	Arf4
0	1	0	2	0	1	0	4	0.0000	0.0001	0.0000	0.0011	Dad1
0	1	2	0	0	17	10	0	0.0000	0.0024	0.0028	0.0000	Sumo2
4	12	8	10	21	84	36	23	0.0162	0.0118	0.0102	0.0065	Hnmpk
0	1	2	3	0	6	8	7	0.0000	0.0008	0.0023	0.0020	Ywhag
0	0	0	1	0	0	0	2	0.0000	0.0000	0.0000	0.0006	Rras2
0	2	0	3	0	4	0	4	0.0000	0.0006	0.0000	0.0011	Rps7
0	1	0	0	0	7	0	0	0.0000	0.0010	0.0000	0.0000	Ppp1ca
0	1	0	0	0	3	0	0	0.0000	0.0004	0.0000	0.0000	Ppp1cb
0	2	1	2	0	4	2	4	0.0000	0.0006	0.0006	0.0011	Calm1
1	2	2	3	2	8	8	8	0.0015	0.0011	0.0023	0.0023	Rps8
0	1	1	2	0	5	2	3	0.0000	0.0007	0.0006	0.0008	Rps15a
3	7	2	3	9	32	7	12	0.0069	0.0045	0.0020	0.0034	Ywhae
1	2	2	1	1	20	10	2	0.0008	0.0028	0.0028	0.0006	Rps14
2	4	4	1	8	22	13	1	0.0062	0.0031	0.0037	0.0003	Rps18
0	0	1	0	0	0	1	0	0.0000	0.0000	0.0003	0.0000	Rps11
0	1	0	0	0	3	0	0	0.0000	0.0004	0.0000	0.0000	Rps13
0	2	0	1	0	4	0	1	0.0000	0.0006	0.0000	0.0003	Snrpe
0	2	1	0	0	8	4	0	0.0000	0.0011	0.0011	0.0000	Snrpf
1	0	0	0	1	0	0	0	0.0008	0.0000	0.0000	0.0000	Snrpg
0	1	0	1	0	1	0	2	0.0000	0.0001	0.0000	0.0006	Snrpd1
0	1	2	0	0	10	3	0	0.0000	0.0014	0.0009	0.0000	Snrpd2
0	1	0	0	0	2	0	0	0.0000	0.0003	0.0000	0.0000	Snrpd3
0	2	3	2	0	3	6	3	0.0000	0.0004	0.0017	0.0008	Rab11a
0	1	1	0	0	2	4	0	0.0000	0.0003	0.0011	0.0000	Tsc22d1
1	3	1	2	1	8	2	6	0.0008	0.0011	0.0006	0.0017	Rps4x
0	0	1	0	0	0	4	0	0.0000	0.0000	0.0011	0.0000	Rpl18a
1	3	4	2	3	8	13	4	0.0023	0.0011	0.0037	0.0011	Rpl23a
0	2	0	0	0	4	0	0	0.0000	0.0006	0.0000	0.0000	Rps6
5	7	5	6	91	206	83	67	0.0702	0.0290	0.0236	0.0189	Hist1h4a
0	0	0	1	0	0	0	2	0.0000	0.0000	0.0000	0.0006	Rab1A
2	3	2	2	8	19	9	7	0.0062	0.0027	0.0026	0.0020	Ran
0	0	0	2	0	0	0	6	0.0000	0.0000	0.0000	0.0017	Rpl23
0	1	1	1	0	11	6	1	0.0000	0.0015	0.0017	0.0003	Rps15
1	0	0	1	1	0	0	2	0.0008	0.0000	0.0000	0.0006	Rps24

0	2	1	2	0	10	7	3	0.0000	0.0014	0.0020	0.0008	Rps25
0	2	1	1	0	10	6	2	0.0000	0.0014	0.0017	0.0006	Rps26
0	2	1	1	0	13	4	3	0.0000	0.0018	0.0011	0.0008	Rps28
0	1	0	0	0	8	0	0	0.0000	0.0011	0.0000	0.0000	Tceb2
0	1	0	1	0	1	0	1	0.0000	0.0001	0.0000	0.0003	Gnb2
0	1	0	2	0	3	0	3	0.0000	0.0004	0.0000	0.0008	Rpl30
1	1	1	1	2	3	3	3	0.0015	0.0004	0.0009	0.0008	Ypl31
1	1	1	3	2	1	7	7	0.0015	0.0001	0.0020	0.0020	Rps3
0	0	1	1	0	0	4	2	0.0000	0.0000	0.0011	0.0006	Rpl32
2	2	2	2	12	10	8	2	0.0093	0.0014	0.0023	0.0006	Rpl8
0	2	1	2	0	11	6	5	0.0000	0.0015	0.0017	0.0014	Ybx1
1	5	1	5	1	19	1	12	0.0008	0.0027	0.0003	0.0034	Pfn1
0	1	0	0	0	1	0	0	0.0000	0.0001	0.0000	0.0000	Crabp1
2	2	2	2	3	19	6	6	0.0023	0.0027	0.0017	0.0017	Uba52
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Tra2b
0	0	1	0	0	0	2	0	0.0000	0.0000	0.0006	0.0000	Pafah1b1
3	19	6	14	13	77	31	38	0.0100	0.0108	0.0088	0.0107	Hspa8
0	0	0	1	0	0	0	2	0.0000	0.0000	0.0000	0.0006	Tpt1
3	9	6	3	18	58	26	8	0.0139	0.0082	0.0074	0.0023	Hspd1
0	1	0	1	0	7	0	1	0.0000	0.0010	0.0000	0.0003	Mapk1
5	11	3	9	14	87	19	29	0.0108	0.0122	0.0054	0.0082	Ywhaz
4	10	7	4	13	36	47	24	0.0100	0.0051	0.0133	0.0068	Hmgb1
1	0	0	0	1	0	0	0	0.0008	0.0000	0.0000	0.0000	Snrpn
1	2	1	4	8	7	2	8	0.0062	0.0010	0.0006	0.0023	Eif5a
0	1	1	1	0	5	6	4	0.0000	0.0007	0.0017	0.0011	Crip1
2	12	5	12	8	99	67	86	0.0062	0.0139	0.0190	0.0242	Actg1
0	1	0	1	0	2	0	1	0.0000	0.0003	0.0000	0.0003	Rps17
0	1	1	1	0	7	2	1	0.0000	0.0010	0.0006	0.0003	Rps12
1	1	1	0	7	2	1	0	0.0054	0.0003	0.0003	0.0000	Rps10
0	0	1	0	0	0	1	0	0.0000	0.0000	0.0003	0.0000	Phb
0	0	1	0	0	0	2	0	0.0000	0.0000	0.0006	0.0000	Csnk2b
0	2	2	2	0	6	16	5	0.0000	0.0008	0.0045	0.0014	Rpl22
0	1	1	0	0	2	2	0	0.0000	0.0003	0.0006	0.0000	Ube2l3
0	0	1	1	0	0	3	2	0.0000	0.0000	0.0009	0.0006	Gnb2l1
3	3	4	5	62	90	42	18	0.0478	0.0127	0.0119	0.0051	Acta1
2	2	1	4	4	21	8	13	0.0031	0.0030	0.0023	0.0037	Ywhaq
1	4	1	4	1	9	9	11	0.0008	0.0013	0.0026	0.0031	Tuba1a
0	0	0	2	0	0	0	4	0.0000	0.0000	0.0000	0.0011	Tubb2c
2	2	1	1	4	3	4	2	0.0031	0.0004	0.0011	0.0006	Hist1h3a
1	3	2	4	1	19	6	5	0.0008	0.0027	0.0017	0.0014	Ywhah
0	1	0	0	0	2	0	0	0.0000	0.0003	0.0000	0.0000	Zhx1
0	0	1	3	0	0	1	5	0.0000	0.0000	0.0003	0.0014	Kpnb1
0	2	1	0	0	3	3	0	0.0000	0.0004	0.0009	0.0000	Psemb7
0	1	0	3	0	7	0	6	0.0000	0.0010	0.0000	0.0017	Pebp1
0	2	1	2	0	10	10	8	0.0000	0.0014	0.0028	0.0023	Hnmph2
0	0	1	1	0	0	8	4	0.0000	0.0000	0.0023	0.0011	Hint1
0	1	1	0	0	9	3	0	0.0000	0.0013	0.0009	0.0000	Elavl1
0	0	1	2	0	0	4	5	0.0000	0.0000	0.0011	0.0014	Slc9a3r1
0	0	1	0	0	0	1	0	0.0000	0.0000	0.0003	0.0000	Cct7
0	1	0	1	0	1	0	1	0.0000	0.0001	0.0000	0.0003	Cct4
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Cct5
0	1	1	1	0	3	3	2	0.0000	0.0004	0.0009	0.0006	Cbx1
0	4	1	4	0	18	2	9	0.0000	0.0025	0.0006	0.0025	Arf1
1	1	1	2	4	6	3	4	0.0031	0.0008	0.0009	0.0011	Rpl19
1	1	0	1	2	2	0	2	0.0015	0.0003	0.0000	0.0006	Srsf3
1	1	0	1	11	2	0	5	0.0085	0.0003	0.0000	0.0014	Hist1h3b
1	1	0	1	2	2	0	2	0.0015	0.0003	0.0000	0.0006	H3f3a
0	0	1	1	0	0	1	1	0.0000	0.0000	0.0003	0.0003	Nptn
0	0	1	0	0	0	1	0	0.0000	0.0000	0.0003	0.0000	Mcm2
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0003	0.0000	Csrp1
0	0	0	1	0	0	0	2	0.0000	0.0000	0.0000	0.0006	Pkp1
0	2	3	1	0	2	4	2	0.0000	0.0003	0.0011	0.0006	Rps3a
0	0	1	1	0	0	2	2	0.0000	0.0000	0.0006	0.0006	S100a13
1	0	1	1	2	0	1	5	0.0015	0.0000	0.0003	0.0014	Psme1
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Psme2
0	1	2	4	0	1	11	9	0.0000	0.0001	0.0031	0.0025	Anxa4
0	0	1	1	0	0	3	1	0.0000	0.0000	0.0009	0.0003	Fhl1
0	0	1	2	0	0	2	3	0.0000	0.0000	0.0006	0.0008	Anpep
0	1	0	0	0	2	0	0	0.0000	0.0003	0.0000	0.0000	Atp5j
0	1	0	0	0	1	0	0	0.0000	0.0001	0.0000	0.0000	Shbg
0	1	0	2	0	2	0	2	0.0000	0.0003	0.0000	0.0006	Fh

0	0	0	2	0	0	0	3	0.0000	0.0000	0.0000	0.0008	G3bp1
0	1	2	8	0	6	9	26	0.0000	0.0008	0.0026	0.0073	Tubb5
0	2	0	0	0	2	0	0	0.0000	0.0003	0.0000	0.0000	Psmb4
3	2	1	2	6	18	3	5	0.0046	0.0025	0.0009	0.0014	Rplp2
0	1	0	3	0	2	0	5	0.0000	0.0003	0.0000	0.0014	Prdx5
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Xdh
0	1	1	3	0	2	1	8	0.0000	0.0003	0.0003	0.0023	Rbp1
0	0	1	0	0	0	4	0	0.0000	0.0000	0.0011	0.0000	Hnmpul2
0	0	0	1	0	0	0	2	0.0000	0.0000	0.0000	0.0006	Rsu1
1	5	4	6	2	28	35	14	0.0015	0.0039	0.0099	0.0039	Nme2
1	4	0	5	1	12	0	11	0.0008	0.0017	0.0000	0.0031	Vcp
1	3	3	8	1	15	9	11	0.0008	0.0021	0.0026	0.0031	Uba1
0	0	0	8	0	0	0	18	0.0000	0.0000	0.0000	0.0051	Jup
1	3	1	5	2	6	10	9	0.0015	0.0008	0.0028	0.0025	Atp5a1
0	0	0	2	0	0	0	2	0.0000	0.0000	0.0000	0.0006	Ckb
0	0	1	1	0	0	2	1	0.0000	0.0000	0.0006	0.0003	Col6a1
0	0	1	1	0	0	4	2	0.0000	0.0000	0.0011	0.0006	Rac2
1	2	1	2	1	6	1	6	0.0008	0.0008	0.0003	0.0017	Rcn1
1	2	2	3	2	13	13	4	0.0015	0.0018	0.0037	0.0011	Fabp5
0	1	1	0	0	2	5	0	0.0000	0.0003	0.0014	0.0000	Atp5i
1	8	4	0	2	16	12	0	0.0015	0.0023	0.0034	0.0000	Igf2r
0	0	1	0	0	0	2	0	0.0000	0.0000	0.0006	0.0000	P311
0	0	0	1	0	0	0	2	0.0000	0.0000	0.0000	0.0000	Tgm3
0	1	0	0	0	2	0	0	0.0000	0.0003	0.0000	0.0000	Dlk1
0	0	0	2	0	0	0	3	0.0000	0.0000	0.0000	0.0008	Krt80
0	0	0	1	0	0	0	4	0.0000	0.0000	0.0000	0.0011	Flg2
0	0	0	1	0	0	0	2	0.0000	0.0000	0.0000	0.0006	Hp1bp3
0	3	3	3	0	14	11	10	0.0000	0.0020	0.0031	0.0028	Myl12b
1	1	0	3	2	1	0	4	0.0015	0.0001	0.0000	0.0011	H2afv
0	1	0	0	0	2	0	0	0.0000	0.0003	0.0000	0.0000	Lig2
1	7	2	5	2	29	9	14	0.0015	0.0041	0.0026	0.0039	Pdia6
0	1	1	0	0	2	4	0	0.0000	0.0003	0.0011	0.0000	Srsf6
1	0	0	0	1	0	0	0	0.0008	0.0000	0.0000	0.0000	Mphosph8
0	3	2	1	0	11	3	1	0.0000	0.0015	0.0009	0.0003	Khsrp
0	0	1	0	0	0	1	0	0.0000	0.0000	0.0003	0.0000	Ddb1
0	0	0	1	0	0	0	2	0.0000	0.0000	0.0000	0.0006	Esy1
0	1	0	0	0	2	0	0	0.0000	0.0003	0.0000	0.0000	Agap2
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Pdap1
1	0	0	0	1	0	0	0	0.0008	0.0000	0.0000	0.0000	Sf3b2
0	1	0	0	0	1	0	0	0.0000	0.0001	0.0000	0.0000	Plcl1
0	0	0	2	0	0	0	3	0.0000	0.0000	0.0000	0.0008	Fam129a
0	0	1	0	0	0	1	0	0.0000	0.0000	0.0003	0.0000	Gm1661
0	0	0	1	0	0	0	2	0.0000	0.0000	0.0000	0.0006	Dnah12
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Lepre1
1	0	0	0	1	0	0	0	0.0008	0.0000	0.0000	0.0000	Psg28
0	0	0	1	0	0	0	2	0.0000	0.0000	0.0000	0.0006	Xirp2
0	1	0	0	0	2	0	0	0.0000	0.0003	0.0000	0.0000	Ppp3ca
1	1	0	2	5	9	0	3	0.0039	0.0013	0.0000	0.0008	Ddx17
0	8	1	0	0	26	3	0	0.0000	0.0037	0.0009	0.0000	Cpa2
0	0	0	1	0	0	0	2	0.0000	0.0000	0.0000	0.0006	DXBay18
0	5	5	7	0	32	18	12	0.0000	0.0045	0.0051	0.0034	Tpm3
1	20	8	12	10	287	55	38	0.0077	0.0404	0.0156	0.0107	Pnliprp1
0	1	0	0	0	1	0	0	0.0000	0.0001	0.0000	0.0000	Rgag4
0	1	0	0	0	1	0	0	0.0000	0.0001	0.0000	0.0000	Trpm3
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Arcn1
0	1	2	3	0	6	16	9	0.0000	0.0008	0.0045	0.0025	Myl6
1	1	0	1	3	16	0	2	0.0023	0.0023	0.0000	0.0006	Akap4
0	2	1	2	0	2	2	5	0.0000	0.0003	0.0006	0.0014	Hnmpd
0	2	1	1	0	16	9	2	0.0000	0.0023	0.0026	0.0006	Psmb6
0	0	0	3	0	0	0	5	0.0000	0.0000	0.0000	0.0014	P4ha1
0	1	1	0	0	6	2	0	0.0000	0.0008	0.0006	0.0000	P4ha2
0	1	1	1	0	1	6	2	0.0000	0.0001	0.0017	0.0006	Bag1
0	1	0	0	0	2	0	0	0.0000	0.0003	0.0000	0.0000	Igf1r
0	2	0	1	0	3	0	2	0.0000	0.0004	0.0000	0.0006	Naca
0	0	0	2	0	0	0	4	0.0000	0.0000	0.0000	0.0011	Serpib6
0	1	1	0	0	1	1	0	0.0000	0.0001	0.0003	0.0000	Stip1
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Caprin1
0	0	1	0	0	0	1	0	0.0000	0.0000	0.0003	0.0000	Reep5
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Vdac3
0	2	3	2	0	13	5	4	0.0000	0.0018	0.0014	0.0011	Vdac1
0	0	2	0	0	0	3	0	0.0000	0.0000	0.0009	0.0000	Dmbt1

0	1	2	2	0	1	5	3	0.0000	0.0001	0.0014	0.0008	Tmpo
0	1	1	0	0	2	1	0	0.0000	0.0003	0.0003	0.0000	Hars
1	5	1	4	3	14	4	7	0.0023	0.0020	0.0011	0.0020	Prdx2
0	1	1	0	0	13	1	0	0.0000	0.0018	0.0003	0.0000	Psap
0	1	0	0	0	2	0	0	0.0000	0.0003	0.0000	0.0000	Pde3b
0	2	0	0	0	6	0	0	0.0000	0.0008	0.0000	0.0000	Hadh
0	2	0	0	0	8	0	0	0.0000	0.0011	0.0000	0.0000	Ewsr1
0	3	3	2	0	5	9	2	0.0000	0.0007	0.0026	0.0006	Fscn1
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Gdi2
0	0	0	1	0	0	0	2	0.0000	0.0000	0.0000	0.0006	Arhgdib
0	1	2	0	0	1	4	0	0.0000	0.0001	0.0011	0.0000	Phgdh
0	0	0	1	0	0	0	2	0.0000	0.0000	0.0000	0.0006	Lasp1
0	0	0	1	0	0	0	2	0.0000	0.0000	0.0000	0.0006	Myh10
2	4	2	4	11	33	15	10	0.0085	0.0046	0.0043	0.0028	Npm1
0	0	0	2	0	0	0	3	0.0000	0.0000	0.0000	0.0008	Pcbp2
0	1	1	1	0	3	6	2	0.0000	0.0004	0.0017	0.0006	Pea15
0	1	0	1	0	9	0	2	0.0000	0.0013	0.0000	0.0006	Srsf2
0	1	0	1	0	1	0	2	0.0000	0.0001	0.0000	0.0006	Aldh1a2
1	0	1	0	1	0	2	0	0.0008	0.0000	0.0006	0.0000	Ddx3x
0	1	1	2	0	2	4	3	0.0000	0.0003	0.0011	0.0008	Dpysl3
0	1	1	0	0	3	1	0	0.0000	0.0004	0.0003	0.0000	Snrpa
0	1	8	27	0	1	23	57	0.0000	0.0001	0.0065	0.0161	Sptbn1
0	1	0	0	0	2	0	0	0.0000	0.0003	0.0000	0.0000	Trim28
0	1	0	0	0	1	0	0	0.0000	0.0001	0.0000	0.0000	Snmp70
0	1	0	0	0	10	0	0	0.0000	0.0014	0.0000	0.0000	Fkbp3
0	1	2	3	0	3	6	6	0.0000	0.0004	0.0017	0.0017	Vat1
0	1	1	1	0	14	6	2	0.0000	0.0020	0.0017	0.0006	Selenbp2
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Sdpr
1	0	1	0	1	0	1	0	0.0008	0.0000	0.0003	0.0000	Raly
0	1	0	0	0	1	0	0	0.0000	0.0001	0.0000	0.0000	Ccpq1
0	6	1	4	0	25	3	9	0.0000	0.0035	0.0009	0.0025	Cel
0	1	0	0	0	2	0	0	0.0000	0.0003	0.0000	0.0000	Cd34
1	3	2	2	10	26	17	4	0.0077	0.0037	0.0048	0.0011	Hspe1
2	2	0	3	30	42	0	14	0.0231	0.0059	0.0000	0.0039	Hist2h2ab
0	1	4	8	0	4	7	14	0.0000	0.0006	0.0020	0.0039	Vcl
0	0	0	2	0	0	0	2	0.0000	0.0000	0.0000	0.0006	Cltc
0	1	0	0	0	3	0	0	0.0000	0.0004	0.0000	0.0000	Morc3
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Nlrp14
1	1	0	1	2	2	0	1	0.0015	0.0003	0.0000	0.0003	Gm9839
1	0	0	3	1	0	0	5	0.0008	0.0000	0.0000	0.0014	Krt77
0	6	4	6	0	34	16	10	0.0000	0.0048	0.0045	0.0028	Tpm4
0	0	1	0	0	0	1	0	0.0000	0.0000	0.0003	0.0000	Sri
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Mapk4
0	1	0	1	0	5	0	1	0.0000	0.0007	0.0000	0.0003	Trak2
0	1	0	0	0	1	0	0	0.0000	0.0001	0.0000	0.0000	Smarcc2
2	4	1	0	7	4	6	0	0.0054	0.0006	0.0017	0.0000	Srsf1
0	0	1	0	0	0	0	1	0.0000	0.0000	0.0003	0.0000	Mylk
0	0	1	0	0	0	2	0	0.0000	0.0000	0.0006	0.0000	Rcor3
0	0	1	0	0	0	1	0	0.0000	0.0000	0.0003	0.0000	Rbm15b
0	1	0	1	0	2	0	2	0.0000	0.0003	0.0000	0.0006	Tpm2
0	1	0	1	0	3	0	2	0.0000	0.0004	0.0000	0.0006	Ubr2
0	2	1	1	0	3	7	2	0.0000	0.0004	0.0020	0.0006	Kctd12
1	1	0	0	1	1	0	0	0.0008	0.0001	0.0000	0.0000	Kdm3b
0	1	1	1	0	6	2	1	0.0000	0.0008	0.0006	0.0003	Rps9
0	0	1	0	0	0	2	0	0.0000	0.0000	0.0006	0.0000	Rps27
0	0	0	2	0	0	0	3	0.0000	0.0000	0.0000	0.0008	Rpl10
1	1	1	1	5	17	3	2	0.0039	0.0024	0.0009	0.0006	Rpl35
0	0	1	0	0	0	1	0	0.0000	0.0000	0.0003	0.0000	Eif2s1
0	3	3	0	0	4	8	0	0.0000	0.0006	0.0023	0.0000	Snd1
0	1	1	0	0	6	7	0	0.0000	0.0008	0.0020	0.0000	Gm5771
0	3	2	0	0	22	6	0	0.0000	0.0031	0.0017	0.0000	Prss3
0	3	1	1	0	7	5	4	0.0000	0.0010	0.0014	0.0011	Try10
0	2	1	3	0	5	6	6	0.0000	0.0007	0.0017	0.0017	Syncrip
0	1	0	0	0	5	0	0	0.0000	0.0007	0.0000	0.0000	Luc7l2
1	0	0	3	1	0	0	5	0.0008	0.0000	0.0000	0.0014	Actn1
1	14	6	7	1	158	39	25	0.0008	0.0222	0.0111	0.0070	Cpa1
1	0	0	0	3	0	0	0	0.0023	0.0000	0.0000	0.0000	Nav3
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Znf579
0	0	1	0	0	0	2	0	0.0000	0.0000	0.0006	0.0000	Mtdh
0	1	1	0	0	2	1	0	0.0000	0.0003	0.0003	0.0000	Jph4
0	0	1	4	0	0	1	7	0.0000	0.0000	0.0003	0.0020	Flnb

0	1	0	0	0	2	0	0	0.0000	0.0003	0.0000	0.0000	BC051142
0	1	0	0	0	3	0	0	0.0000	0.0004	0.0000	0.0000	Nucks1
0	0	1	0	0	0	2	0	0.0000	0.0000	0.0000	0.0006	Tufm
0	0	1	0	0	0	1	0	0.0000	0.0000	0.0003	0.0000	Cpped1
2	4	4	6	24	36	21	20	0.0185	0.0051	0.0060	0.0056	Hnrmpa3
1	0	0	0	1	0	0	0	0.0008	0.0000	0.0000	0.0000	Eif4b
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Mak16
0	0	0	3	0	0	0	7	0.0000	0.0000	0.0000	0.0020	Ehd2
0	0	1	0	0	0	3	0	0.0000	0.0000	0.0009	0.0000	Echs1
0	2	1	4	0	4	3	7	0.0000	0.0006	0.0009	0.0020	Rcn3
0	1	0	0	0	1	0	0	0.0000	0.0001	0.0000	0.0000	Ganab
0	1	0	0	0	1	0	0	0.0000	0.0001	0.0000	0.0000	Sgta
0	1	0	0	0	1	0	0	0.0000	0.0001	0.0000	0.0000	Rcc2
0	2	1	0	0	4	2	0	0.0000	0.0006	0.0006	0.0000	Srsf7
2	8	6	9	6	38	22	23	0.0046	0.0053	0.0062	0.0065	Ckap4
0	0	3	2	0	0	5	4	0.0000	0.0000	0.0014	0.0011	Hadha
0	0	1	1	0	0	1	2	0.0000	0.0000	0.0003	0.0006	Rpl24
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Rcn2
0	0	0	1	0	0	0	2	0.0000	0.0000	0.0000	0.0000	Asph
0	0	1	0	0	0	1	0	0.0000	0.0000	0.0003	0.0000	Lig4
0	10	12	6	0	30	36	9	0.0000	0.0042	0.0102	0.0025	Flna
0	0	1	1	0	0	2	2	0.0000	0.0000	0.0006	0.0006	Aifm2
0	0	0	4	0	0	0	6	0.0000	0.0000	0.0000	0.0017	Acaa2
0	1	0	0	0	1	0	0	0.0000	0.0001	0.0000	0.0000	Pxk
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Asrgl1
0	0	0	1	0	0	0	2	0.0000	0.0000	0.0000	0.0006	ncharacterized
0	3	1	1	0	10	2	1	0.0000	0.0014	0.0006	0.0003	11-Sep
0	0	1	0	0	0	1	0	0.0000	0.0000	0.0003	0.0000	Gykl1
1	0	0	0	1	0	0	0	0.0008	0.0000	0.0000	0.0000	Tceal5
0	1	2	0	0	3	6	0	0.0000	0.0004	0.0017	0.0000	Txn1
0	1	0	0	0	1	0	0	0.0000	0.0001	0.0000	0.0000	Qrfp
4	6	5	6	117	171	37	52	0.0903	0.0241	0.0105	0.0146	Hist1h2bp
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Copa
0	0	0	1	0	0	0	2	0.0000	0.0000	0.0000	0.0006	Fgb
0	1	1	1	0	6	2	2	0.0000	0.0008	0.0006	0.0006	Tpm3
1	1	1	1	3	3	6	2	0.0023	0.0004	0.0017	0.0006	Sdha
1	1	0	0	1	1	0	0	0.0008	0.0001	0.0000	0.0000	P2rx2
0	0	1	0	0	0	2	0	0.0000	0.0000	0.0006	0.0000	Pof1b
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Lgi2
0	2	0	3	0	2	0	6	0.0000	0.0003	0.0000	0.0017	Acat1
0	1	0	0	0	1	0	0	0.0000	0.0001	0.0000	0.0000	Gspt1
0	4	0	0	0	7	0	0	0.0000	0.0010	0.0000	0.0000	Hnrmp1
1	1	0	1	11	36	0	16	0.0085	0.0051	0.0000	0.0045	H2afj
0	1	1	1	0	3	6	6	0.0000	0.0004	0.0017	0.0017	Ssspta
0	0	1	0	0	0	2	0	0.0000	0.0000	0.0006	0.0000	Ushbp1
0	1	1	0	0	1	2	0	0.0000	0.0001	0.0006	0.0000	Actr1b
1	0	0	0	1	0	0	0	0.0008	0.0000	0.0000	0.0000	Sgms1
0	3	1	3	0	9	4	6	0.0000	0.0013	0.0011	0.0017	Cald1
0	0	3	15	0	0	7	26	0.0000	0.0000	0.0020	0.0073	Myh9
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Atp1a1
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	rotein
1	1	1	1	1	4	5	1	0.0008	0.0006	0.0014	0.0003	Srsf4
1	3	3	5	2	15	11	9	0.0015	0.0021	0.0031	0.0025	Hnrmpu
0	0	1	0	0	0	1	0	0.0000	0.0000	0.0003	0.0000	Olf1317
0	1	0	0	0	1	0	0	0.0000	0.0001	0.0000	0.0000	Olf1516
0	2	1	2	0	3	2	2	0.0000	0.0004	0.0006	0.0006	Hnrmp1
1	3	3	1	4	18	14	5	0.0031	0.0025	0.0040	0.0014	Sfpq
0	1	0	0	0	2	0	0	0.0000	0.0003	0.0000	0.0000	Slc12a5
0	1	0	0	0	2	0	0	0.0000	0.0003	0.0000	0.0000	Rab14
0	0	0	1	0	0	0	3	0.0000	0.0000	0.0000	0.0008	Rps5
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Isoc1
1	1	2	3	3	16	13	16	0.0023	0.0023	0.0037	0.0045	Hba-a1
0	0	4	1	0	0	7	2	0.0000	0.0000	0.0020	0.0006	Rnh1
0	1	1	2	0	1	2	2	0.0000	0.0001	0.0006	0.0006	Rbmx11
0	0	1	0	0	0	1	0	0.0000	0.0000	0.0003	0.0000	Ppa2
0	3	1	3	0	6	6	9	0.0000	0.0008	0.0017	0.0025	Txndc5
0	1	0	1	0	4	0	1	0.0000	0.0006	0.0000	0.0003	Fubp1
0	0	1	0	0	0	1	0	0.0000	0.0000	0.0003	0.0000	Yars
0	4	1	4	0	28	4	11	0.0000	0.0039	0.0011	0.0031	Cela1
3	8	6	4	6	82	27	6	0.0046	0.0115	0.0077	0.0017	Basp1
0	1	0	0	0	2	0	0	0.0000	0.0003	0.0000	0.0000	F11

0	0	1	0	0	0	1	0	0.0000	0.0000	0.0003	0.0000	Wasl
0	1	1	3	0	1	4	6	0.0000	0.0001	0.0011	0.0017	Rpn1
1	1	1	1	7	3	1	1	0.0054	0.0004	0.0003	0.0003	Nrif2
0	1	0	1	0	3	0	3	0.0000	0.0004	0.0000	0.0008	Tardbp
0	1	0	0	0	2	0	0	0.0000	0.0003	0.0000	0.0000	Tf
0	0	0	1	0	0	0	2	0.0000	0.0000	0.0000	0.0006	Tubb6
1	3	2	1	1	8	3	2	0.0008	0.0011	0.0009	0.0006	Ptbp1
1	4	2	5	6	22	14	24	0.0046	0.0031	0.0040	0.0068	Krt5
0	1	0	0	0	4	0	0	0.0000	0.0006	0.0000	0.0000	Sf3b5
0	1	0	3	0	2	0	4	0.0000	0.0003	0.0000	0.0011	Taldo1
0	2	0	2	0	17	0	6	0.0000	0.0024	0.0000	0.0017	Hnmpab
0	0	0	1	0	0	0	2	0.0000	0.0000	0.0000	0.0006	Nans
0	0	1	0	0	0	1	0	0.0000	0.0000	0.0003	0.0000	Psip1
0	1	1	1	0	4	3	2	0.0000	0.0006	0.0009	0.0006	Rap1b
0	1	0	0	0	1	0	0	0.0000	0.0001	0.0000	0.0000	Ggt7
0	0	0	3	0	0	0	4	0.0000	0.0000	0.0000	0.0011	Actr3
0	2	1	0	0	4	4	0	0.0000	0.0006	0.0011	0.0000	Nono
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Vwa5a
0	0	1	0	0	0	2	0	0.0000	0.0000	0.0006	0.0000	Card14
0	2	1	2	0	4	2	5	0.0000	0.0006	0.0006	0.0014	Tmed9
0	0	1	1	0	0	2	2	0.0000	0.0000	0.0006	0.0006	Aco2
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Dctn2
0	0	0	1	0	0	0	2	0.0000	0.0000	0.0000	0.0006	Dnajb11
0	0	0	1	0	0	0	2	0.0000	0.0000	0.0000	0.0006	St13
0	1	0	1	0	2	0	3	0.0000	0.0003	0.0000	0.0008	Etfa
0	0	1	0	0	0	2	0	0.0000	0.0000	0.0006	0.0000	Ddah2
0	1	0	0	0	1	0	0	0.0000	0.0001	0.0000	0.0000	Gpx7
0	0	1	2	0	0	1	3	0.0000	0.0000	0.0003	0.0008	Park7
0	1	1	0	0	10	3	0	0.0000	0.0014	0.0009	0.0000	Krt82
0	2	1	0	0	9	2	0	0.0000	0.0013	0.0006	0.0000	Nasp
1	0	0	0	1	0	0	0	0.0008	0.0000	0.0000	0.0000	Srrt
0	1	0	3	0	1	0	4	0.0000	0.0001	0.0000	0.0011	Rtn4
0	8	6	11	0	34	20	20	0.0000	0.0048	0.0057	0.0056	Rrbp1
0	0	0	1	0	0	0	3	0.0000	0.0000	0.0000	0.0008	Krt23
1	2	0	4	2	5	0	8	0.0015	0.0007	0.0000	0.0023	Arhgdia
0	1	1	2	0	2	4	3	0.0000	0.0003	0.0011	0.0008	2210010C04Rik
1	1	0	0	2	2	0	0	0.0015	0.0003	0.0000	0.0000	Cox6c
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Cenpq
0	1	2	3	0	5	11	7	0.0000	0.0007	0.0031	0.0020	Rpl17
0	0	1	0	0	0	2	0	0.0000	0.0000	0.0006	0.0000	Glod4
0	0	1	0	0	0	2	0	0.0000	0.0000	0.0006	0.0000	Arpc5
0	5	2	5	0	61	14	11	0.0000	0.0086	0.0040	0.0031	Cela3b
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Mtap
0	4	0	0	0	18	0	0	0.0000	0.0025	0.0000	0.0000	Clps
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Pcyox1
0	1	0	0	0	1	0	0	0.0000	0.0001	0.0000	0.0000	Cott1
0	0	1	0	0	0	1	0	0.0000	0.0000	0.0003	0.0000	Atp5f1
0	2	3	2	0	16	13	2	0.0000	0.0023	0.0037	0.0006	Rps21
1	5	3	7	2	41	12	17	0.0015	0.0058	0.0034	0.0048	Ywhab
2	8	2	4	21	233	9	14	0.0162	0.0328	0.0026	0.0039	Ctrb1
0	2	2	3	0	15	11	8	0.0000	0.0021	0.0031	0.0023	Rpl14
0	1	2	7	0	2	4	11	0.0000	0.0003	0.0011	0.0031	Tppp3
1	1	0	0	2	1	0	0	0.0015	0.0001	0.0000	0.0000	Filip1
0	3	1	5	0	6	11	14	0.0000	0.0008	0.0031	0.0039	Tubb2b
0	1	0	0	0	2	0	0	0.0000	0.0003	0.0000	0.0000	Rbm8a
0	0	0	1	0	0	0	2	0.0000	0.0000	0.0000	0.0006	Cygb
0	0	0	1	0	0	0	2	0.0000	0.0000	0.0000	0.0006	Hnmpa0
0	1	0	1	0	1	0	1	0.0000	0.0001	0.0000	0.0003	Rpl11
1	2	1	2	2	5	11	4	0.0015	0.0007	0.0031	0.0011	Ssr1
1	2	0	2	1	8	0	4	0.0008	0.0011	0.0000	0.0011	Serbp1
0	0	1	1	0	0	1	1	0.0000	0.0000	0.0003	0.0003	Tpd52l2
0	0	1	0	0	0	1	0	0.0000	0.0000	0.0003	0.0000	Uqcrc1
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Cmtm6
0	0	1	0	0	0	1	0	0.0000	0.0000	0.0003	0.0000	Rpl15
0	1	0	0	0	1	0	0	0.0000	0.0001	0.0000	0.0000	Shmt2
2	4	1	3	8	19	4	4	0.0062	0.0027	0.0011	0.0011	Rps19
0	0	1	0	0	0	1	0	0.0000	0.0000	0.0003	0.0000	Pdhb
0	1	1	1	0	1	3	2	0.0000	0.0001	0.0009	0.0006	Hnmpm
0	1	1	0	0	1	1	0	0.0000	0.0001	0.0003	0.0000	Pgm1
1	1	1	1	2	15	9	5	0.0015	0.0021	0.0026	0.0014	Ptms
0	1	0	0	0	2	0	0	0.0000	0.0003	0.0000	0.0000	Oxct1

0	0	1	0	0	0	1	0	0.0000	0.0000	0.0003	0.0000	Nhp211
0	2	0	3	0	4	0	4	0.0000	0.0006	0.0000	0.0011	Tmed10
0	1	0	0	0	2	0	0	0.0000	0.0003	0.0000	0.0000	Cthrc1
0	1	1	0	0	1	2	0	0.0000	0.0001	0.0006	0.0000	Sarp
0	0	1	0	0	0	2	0	0.0000	0.0000	0.0006	0.0000	Sec13
0	1	1	1	0	5	3	1	0.0000	0.0007	0.0009	0.0003	Rpl34
0	0	0	1	0	0	0	2	0.0000	0.0000	0.0000	0.0006	Fam114a1
0	1	1	1	0	2	6	2	0.0000	0.0003	0.0017	0.0006	Ube2v2
0	1	0	0	0	1	0	0	0.0000	0.0001	0.0000	0.0000	Acp1
1	2	1	0	9	8	1	0	0.0069	0.0011	0.0003	0.0000	Atp5d
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Sec23b
0	1	0	0	0	1	0	0	0.0000	0.0001	0.0000	0.0000	Drap1
0	0	1	1	0	0	3	1	0.0000	0.0000	0.0009	0.0003	Mrpl2
0	1	1	1	0	3	5	1	0.0000	0.0004	0.0014	0.0003	Chmp4b
0	0	1	1	0	0	4	2	0.0000	0.0000	0.0011	0.0000	Rpl4
0	1	3	1	0	2	10	2	0.0000	0.0003	0.0028	0.0006	Eef1g
0	1	1	0	0	4	2	0	0.0000	0.0006	0.0006	0.0000	C1qtnf2
0	1	0	0	0	1	0	0	0.0000	0.0001	0.0000	0.0000	Phpt1
0	2	0	0	0	5	0	0	0.0000	0.0007	0.0000	0.0000	Cnn3
1	1	1	0	1	2	1	0	0.0008	0.0003	0.0003	0.0000	Uqcrc2
0	0	1	0	0	0	1	0	0.0000	0.0000	0.0003	0.0000	Plin3
1	0	0	1	3	0	0	2	0.0023	0.0000	0.0000	0.0006	Rpn2
2	5	3	3	5	23	6	6	0.0039	0.0032	0.0017	0.0017	Pgam1
0	0	1	0	0	0	1	0	0.0000	0.0000	0.0003	0.0000	Cmpk1
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Ikbip
0	1	0	0	0	2	0	0	0.0000	0.0003	0.0000	0.0000	Abcb6
0	0	1	1	0	0	5	1	0.0000	0.0000	0.0014	0.0003	Pgd
0	0	0	1	0	0	0	2	0.0000	0.0000	0.0000	0.0006	Ssr3
0	0	0	1	0	0	0	2	0.0000	0.0000	0.0000	0.0006	Paics
0	1	3	4	0	2	13	8	0.0000	0.0003	0.0037	0.0023	Cyb5r3
0	1	1	0	0	2	1	0	0.0000	0.0003	0.0003	0.0000	Aspdh
0	0	0	3	0	0	0	6	0.0000	0.0000	0.0000	0.0017	Krt7
0	2	3	2	0	5	8	2	0.0000	0.0007	0.0023	0.0006	Etfb
0	0	1	0	0	0	8	0	0.0000	0.0000	0.0023	0.0000	Atp5h
0	1	0	0	0	2	0	0	0.0000	0.0003	0.0000	0.0000	Fgf23
1	3	1	3	4	19	2	5	0.0031	0.0027	0.0006	0.0014	Set
0	1	1	5	0	4	4	10	0.0000	0.0006	0.0011	0.0028	Ctrl
0	0	1	2	0	0	1	5	0.0000	0.0000	0.0003	0.0014	Gsdma
0	1	1	0	0	2	4	0	0.0000	0.0003	0.0011	0.0000	Anp32b
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Tmod3
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Dync1h1
1	0	0	0	1	0	0	0	0.0008	0.0000	0.0000	0.0000	Isyna1
0	2	1	1	0	3	2	1	0.0000	0.0004	0.0006	0.0003	Akr1a1
0	1	0	0	0	1	0	0	0.0000	0.0001	0.0000	0.0000	Ralb
0	1	0	0	0	1	0	0	0.0000	0.0001	0.0000	0.0000	Caln1
0	1	1	1	0	9	3	1	0.0000	0.0013	0.0009	0.0003	Sh3bgr1
0	1	1	1	0	2	6	1	0.0000	0.0003	0.0017	0.0003	Refbp2
0	1	0	0	0	1	0	0	0.0000	0.0001	0.0000	0.0000	Myg1
0	1	0	0	0	2	0	0	0.0000	0.0003	0.0000	0.0000	Serpini2
0	1	0	0	0	3	0	0	0.0000	0.0004	0.0000	0.0000	Aldh9a1
0	0	1	0	0	0	1	0	0.0000	0.0000	0.0003	0.0000	Klra17
0	0	1	0	0	0	1	0	0.0000	0.0000	0.0003	0.0000	Hdgfrp3
0	1	2	1	0	1	7	2	0.0000	0.0001	0.0020	0.0006	Rhoa
0	0	1	1	0	0	6	3	0.0000	0.0000	0.0017	0.0008	Psma6
0	0	1	0	0	0	1	0	0.0000	0.0000	0.0003	0.0000	Kif21b
0	0	1	0	0	0	1	0	0.0000	0.0000	0.0003	0.0000	Plec
0	1	0	1	0	3	0	2	0.0000	0.0004	0.0000	0.0006	Cnpy2
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Clic4
0	0	0	3	0	0	0	5	0.0000	0.0000	0.0000	0.0014	Add3
0	0	2	2	0	0	5	4	0.0000	0.0000	0.0014	0.0011	Add1
0	1	0	0	0	1	0	0	0.0000	0.0001	0.0000	0.0000	Eif3i
0	1	0	0	0	1	0	0	0.0000	0.0001	0.0000	0.0000	Galk1
0	0	1	0	0	0	1	0	0.0000	0.0000	0.0003	0.0000	Esd
0	0	0	4	0	0	0	7	0.0000	0.0000	0.0000	0.0020	Dstn
0	1	0	0	0	1	0	0	0.0000	0.0001	0.0000	0.0000	Sec11a
1	2	1	1	1	7	2	2	0.0008	0.0010	0.0006	0.0006	Tmed2
0	0	0	2	0	0	0	3	0.0000	0.0000	0.0000	0.0008	Try4
0	1	1	1	0	5	3	1	0.0000	0.0007	0.0009	0.0003	Ikbke
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Ak1
0	2	0	0	0	4	0	0	0.0000	0.0006	0.0000	0.0000	Psma4
0	1	1	1	0	4	3	2	0.0000	0.0006	0.0009	0.0006	Psmb2

0	2	1	0	0	5	1	0	0.0000	0.0007	0.0003	0.0000	Psma1
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Arntl
0	2	0	0	0	7	0	0	0.0000	0.0010	0.0000	0.0000	Ak2
0	8	9	10	0	52	36	14	0.0000	0.0073	0.0102	0.0039	Akap12
0	1	1	0	0	5	4	0	0.0000	0.0007	0.0011	0.0000	Skp1
0	0	0	1	0	0	0	2	0.0000	0.0000	0.0000	0.0006	Eif4h
0	0	0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Coro1c
0	1	0	0	0	3	0	0	0.0000	0.0004	0.0000	0.0000	Ctsz
0	0	1	1	0	0	5	1	0.0000	0.0000	0.0014	0.0003	Snx1
0	2	2	5	0	10	7	15	0.0000	0.0014	0.0020	0.0042	Tagln2
0	0	0	2	0	0	0	3	0.0000	0.0000	0.0000	0.0008	Ehd1
0	0	0	2	0	0	0	3	0.0000	0.0000	0.0000	0.0008	Gbp2
0	2	0	1	0	2	0	1	0.0000	0.0003	0.0000	0.0003	Aifm1
1	1	0	0	1	2	0	0	0.0008	0.0003	0.0000	0.0000	Ddx39b
0	2	1	1	0	3	4	1	0.0000	0.0004	0.0011	0.0003	Clic1
0	1	1	0	0	3	3	0	0.0000	0.0004	0.0009	0.0000	Vars
0	1	0	1	0	2	0	5	0.0000	0.0003	0.0000	0.0014	Prss1
0	1	0	0	0	2	0	0	0.0000	0.0003	0.0000	0.0000	Thbs4
0	1	0	0	0	2	0	0	0.0000	0.0003	0.0000	0.0000	Strap
0	4	0	2	0	6	0	2	0.0000	0.0008	0.0000	0.0006	Hnmpc
0	0	1	0	0	0	2	0	0.0000	0.0000	0.0006	0.0000	Pmm2
0	2	2	2	0	6	9	6	0.0000	0.0008	0.0026	0.0017	Psma7
0	0	0	1	0	0	0	2	0.0000	0.0000	0.0000	0.0006	Psma5
0	0	1	0	0	0	1	0	0.0000	0.0000	0.0003	0.0000	Dnpep
0	1	0	1	0	4	0	1	0.0000	0.0006	0.0000	0.0003	Hnmpf

protein	annotation	FDR				Unique Petides			
		e15.5	e17.5	p2	p14	e15.5	e17.5	p2	
E9Q447	Uncharacterized protein OS=Mus musculus GN=Spna2 PE=4 SV=1	1.00	0.00	0.00	0.00	0	3	8	
O08547	Vesicle-trafficking protein SEC22b OS=Mus musculus GN=Sec22b PE=1 SV=3	1.00	1.00	1.00	0.02	0	0	0	
O35074	Prostacyclin synthase OS=Mus musculus GN=Ptgis PE=2 SV=1	1.00	1.00	1.00	0.00	0	0	0	
O35129	Prohibitin-2 OS=Mus musculus GN=Phb2 PE=1 SV=1	1.00	0.02	0.02	1.00	0	1	1	
O35682	Myeloid-associated differentiation marker OS=Mus musculus GN=Myadm PE=2 SV=2	1.00	1.00	1.00	0.02	0	0	0	
O54734	Dolichyl-diphosphooligosaccharide--protein glycosyltransferase 48 kDa subunit OS=Mus musculus	1.00	0.02	1.00	0.00	0	1	0	
O55022	Membrane-associated progesterone receptor component 1 OS=Mus musculus GN=Pgrmc1 PE=	1.00	0.02	1.00	1.00	0	1	0	
O55026	Ectonucleoside triphosphate diphosphohydrolase 2 OS=Mus musculus GN=Entpd2 PE=1 SV=2	1.00	1.00	1.00	0.00	0	0	0	
O55029	Coatomer subunit beta' OS=Mus musculus GN=Copb2 PE=2 SV=2	1.00	1.00	1.00	0.02	0	0	0	
O88962	7-alpha-hydroxycholest-4-en-3-one 12-alpha-hydroxylase OS=Mus musculus GN=Cyp8b1 PE=2	1.00	1.00	1.00	0.07	0	0	0	
P05064	Fructose-bisphosphate aldolase A OS=Mus musculus GN=Aldoa PE=1 SV=2	0.02	0.00	0.00	0.00	1	9	4	
P11688	Integrin alpha-5 OS=Mus musculus GN=Itga5 PE=1 SV=3	1.00	0.04	0.02	1.00	0	1	1	
P14431	H-2 class I histocompatibility antigen, Q9 alpha chain (Fragment) OS=Mus musculus GN=H2-Q9	1.00	0.04	1.00	1.00	0	1	0	
P14733	Lamin-B1 OS=Mus musculus GN=Lmnb1 PE=1 SV=3	0.00	0.00	0.00	0.00	2	5	5	
P21981	Protein-glutamine gamma-glutamyltransferase 2 OS=Mus musculus GN=Tgm2 PE=1 SV=4	1.00	1.00	1.00	0.00	0	0	0	
P26645	Myristoylated alanine-rich C-kinase substrate OS=Mus musculus GN=Marcks PE=1 SV=2	0.00	0.00	0.00	0.00	5	6	4	
P35564	Calnexin OS=Mus musculus GN=Canx PE=1 SV=1	1.00	0.00	1.00	0.00	0	2	0	
P45878	Peptidyl-prolyl cis-trans isomerase FKBP2 OS=Mus musculus GN=Fkbp2 PE=1 SV=1	1.00	0.02	1.00	1.00	0	1	0	
P47740	Fatty aldehyde dehydrogenase OS=Mus musculus GN=Aldh3a2 PE=2 SV=2	1.00	1.00	1.00	0.02	0	0	0	
P47753	F-actin-capping protein subunit alpha-1 OS=Mus musculus GN=Capza1 PE=1 SV=4	1.00	0.02	0.02	0.02	0	1	1	
P47757	F-actin-capping protein subunit beta OS=Mus musculus GN=Capzb PE=1 SV=3	1.00	0.03	0.03	0.02	0	1	1	
P48771	Cytochrome c oxidase subunit 7A2, mitochondrial OS=Mus musculus GN=Cox7a2 PE=1 SV=2	1.00	0.02	1.00	1.00	0	1	0	
P56395	Cytochrome b5 OS=Mus musculus GN=Cyb5a PE=1 SV=2	1.00	1.00	1.00	0.02	0	0	0	
P56501	Mitochondrial uncoupling protein 3 OS=Mus musculus GN=Ucp3 PE=2 SV=1	0.05	0.03	1.00	0.08	1	1	0	
P61168	D(2) dopamine receptor OS=Mus musculus GN=Drd2 PE=2 SV=2	0.04	0.03	1.00	1.00	1	1	0	
P61804	Dolichyl-diphosphooligosaccharide--protein glycosyltransferase subunit DAD1 OS=Mus musculus	1.00	0.03	1.00	0.00	0	1	0	
P63005	Platelet-activating factor acetylhydrolase IB subunit alpha OS=Mus musculus GN=Pafah1b1 PE=	1.00	1.00	0.02	1.00	0	0	1	
P63242	Eukaryotic translation initiation factor 5A-1 OS=Mus musculus GN=Elf5a PE=1 SV=2	0.02	0.00	0.04	0.00	1	2	1	
P67778	Prohibitin OS=Mus musculus GN=Phb PE=1 SV=1	1.00	1.00	0.02	1.00	0	0	1	
P70441	Na(+)/H(+) exchange regulatory cofactor NHE-RF1 OS=Mus musculus GN=Slc9a3r1 PE=1 SV=3	1.00	1.00	0.02	0.00	0	0	1	
P97449	Aminopeptidase N OS=Mus musculus GN=Anpep PE=1 SV=4	1.00	1.00	0.03	0.00	0	0	1	
P97450	ATP synthase-coupling factor 6, mitochondrial OS=Mus musculus GN=Atp5j PE=1 SV=1	1.00	0.04	1.00	1.00	0	1	0	
Q02257	Junction plakoglobin OS=Mus musculus GN=Jup PE=1 SV=3	1.00	1.00	1.00	0.00	0	0	0	
Q05144	Ras-related C3 botulinum toxin substrate 2 OS=Mus musculus GN=Rac2 PE=2 SV=1	1.00	1.00	0.02	0.02	0	0	1	
Q06185	ATP synthase subunit e, mitochondrial OS=Mus musculus GN=Atp5i PE=1 SV=2	1.00	0.02	0.02	1.00	0	1	1	
Q07113	Cation-independent mannose-6-phosphate receptor OS=Mus musculus GN=Igf2r PE=1 SV=1	0.02	0.00	0.00	1.00	1	8	4	
Q09163	Protein delta homolog 1 OS=Mus musculus GN=Dlk1 PE=1 SV=1	1.00	0.04	1.00	1.00	0	1	0	
Q3U7R1	Extended synaptotagmin-1 OS=Mus musculus GN=Esyt1 PE=2 SV=2	1.00	1.00	1.00	0.03	0	0	0	
Q5F4T0	Transient receptor potential cation channel, subfamily M, member 3 OS=Mus musculus GN=Trpr	1.00	0.02	1.00	1.00	0	1	0	
Q5XJY5	Coatomer subunit delta OS=Mus musculus GN=Arcn1 PE=2 SV=2	1.00	1.00	1.00	0.02	0	0	0	
Q60870	Receptor expression-enhancing protein 5 OS=Mus musculus GN=Reep5 PE=1 SV=1	1.00	1.00	0.02	1.00	0	0	1	
Q60931	Voltage-dependent anion-selective channel protein 3 OS=Mus musculus GN=Vdac3 PE=1 SV=1	1.00	1.00	1.00	0.07	0	0	0	
Q60932	Voltage-dependent anion-selective channel protein 1 OS=Mus musculus GN=Vdac1 PE=1 SV=3	1.00	0.00	0.00	0.00	0	2	3	
Q60997	Deleted in malignant brain tumors 1 protein OS=Mus musculus GN=Dmbt1 PE=1 SV=2	1.00	1.00	0.00	1.00	0	0	2	
Q61029	Lamina-associated polypeptide 2, isoforms beta/delta/epsilon/gamma OS=Mus musculus GN=Trr	1.00	0.02	0.00	0.00	0	1	2	
Q61409	cGMP-inhibited 3',5'-cyclic phosphodiesterase B OS=Mus musculus GN=Pde3b PE=1 SV=2	1.00	0.02	1.00	1.00	0	1	0	
Q61598	Rab GDP dissociation inhibitor beta OS=Mus musculus GN=Gdi2 PE=1 SV=1	1.00	1.00	1.00	0.02	0	0	0	
Q62261	Spectrin beta chain, brain 1 OS=Mus musculus GN=Sptbn1 PE=1 SV=2	1.00	0.05	0.00	0.00	0	1	8	
Q63836	Selenium-binding protein 2 OS=Mus musculus GN=Selenbp2 PE=1 SV=2	1.00	0.02	0.02	0.02	0	1	1	
Q640L3	Cell cycle progression protein 1 OS=Mus musculus GN=Ccpg1 PE=1 SV=2	1.00	0.04	1.00	1.00	0	1	0	
Q64314	Hematopoietic progenitor cell antigen CD34 OS=Mus musculus GN=Cd34 PE=1 SV=1	1.00	0.07	1.00	1.00	0	1	0	
Q68FD5	Clathrin heavy chain 1 OS=Mus musculus GN=Cltc PE=1 SV=3	1.00	1.00	1.00	0.00	0	0	0	
Q80TN7	Neuron navigator 3 OS=Mus musculus GN=Nav3 PE=1 SV=2	0.04	1.00	1.00	1.00	1	0	0	
Q80WJ7	Protein LYRIC OS=Mus musculus GN=Mtdh PE=1 SV=1	1.00	1.00	0.05	1.00	0	0	1	
Q8BH64	EH domain-containing protein 2 OS=Mus musculus GN=Ehd2 PE=1 SV=1	1.00	1.00	1.00	0.00	0	0	0	
Q8BSY0	Aspartyl/asparaginyl beta-hydroxylase OS=Mus musculus GN=Asph PE=2 SV=1	1.00	1.00	1.00	0.02	0	0	0	
Q8BUE4	Apoptosis-inducing factor 2 OS=Mus musculus GN=Aifm2 PE=2 SV=1	1.00	1.00	0.05	0.02	0	0	1	
Q8CIE6	Coatomer subunit alpha OS=Mus musculus GN=Copa PE=1 SV=2	1.00	1.00	1.00	0.02	0	0	0	
Q8K2B3	Succinate dehydrogenase [ubiquinone] flavoprotein subunit, mitochondrial OS=Mus musculus GN	0.03	0.02	0.02	0.02	1	1	1	
Q8R207	Small subunit of serine palmitoyltransferase A OS=Mus musculus GN=Ssspta PE=2 SV=2	1.00	0.02	0.02	0.02	0	1	1	
Q91YQ5	Dolichyl-diphosphooligosaccharide--protein glycosyltransferase subunit 1 OS=Mus musculus GN	1.00	0.04	0.03	0.00	0	1	1	
Q99JP7	Gamma-glutamyltransferase 7 OS=Mus musculus GN=Ggt7 PE=1 SV=2	1.00	0.07	1.00	1.00	0	1	0	
Q99KF1	Transmembrane emp24 domain-containing protein 9 OS=Mus musculus GN=Tmed9 PE=2 SV=2	1.00	0.00	0.02	0.00	0	2	1	
Q99KJ8	Dynactin subunit 2 OS=Mus musculus GN=Dctn2 PE=1 SV=3	1.00	1.00	1.00	0.02	0	0	0	
Q99PL5	Ribosome-binding protein 1 OS=Mus musculus GN=Rrbp1 PE=2 SV=2	1.00	0.00	0.00	0.00	0	8	6	
Q9CPQ1	Cytochrome c oxidase subunit 6C OS=Mus musculus GN=Cox6c PE=1 SV=3	0.04	0.02	1.00	1.00	1	1	0	
Q9CQQ7	ATP synthase subunit b, mitochondrial OS=Mus musculus GN=Atp5f1 PE=1 SV=1	1.00	1.00	0.05	1.00	0	0	1	
Q9CY50	Translocon-associated protein subunit alpha OS=Mus musculus GN=Ssr1 PE=1 SV=1	0.06	0.00	0.02	0.00	1	2	1	
Q9CZ13	Cytochrome b-c1 complex subunit 1, mitochondrial OS=Mus musculus GN=Uqcrc1 PE=1 SV=2	1.00	1.00	0.02	1.00	0	0	1	
Q9CZ69	CKLF-like MARVEL transmembrane domain-containing protein 6 OS=Mus musculus GN=Cmtm6	1.00	1.00	1.00	0.04	0	0	0	
Q9D1M0	Protein SEC13 homolog OS=Mus musculus GN=Sec13 PE=2 SV=3	1.00	1.00	0.02	1.00	0	0	1	
Q9D3D9	ATP synthase subunit delta, mitochondrial OS=Mus musculus GN=Atp5d PE=1 SV=1	0.02	0.00	0.03	1.00	1	2	1	
Q9D662	Protein transport protein Sec23B OS=Mus musculus GN=Sec23b PE=2 SV=1	1.00	1.00	1.00	0.02	0	0	0	

Q9D8B3	Charged multivesicular body protein 4b OS=Mus musculus GN=Chmp4b PE=2 SV=2	1.00	0.02	0.02	0.02	0	1	1
Q9DB77	Cytochrome b-c1 complex subunit 2, mitochondrial OS=Mus musculus GN=Uqcrc2 PE=1 SV=1	0.04	0.02	0.02	1.00	1	1	1
Q9DBG5	Perilipin-3 OS=Mus musculus GN=Plin3 PE=1 SV=1	1.00	1.00	0.03	1.00	0	0	1
Q9DBG6	Dolichyl-diphosphooligosaccharide--protein glycosyltransferase subunit 2 OS=Mus musculus GN:	0.03	1.00	1.00	0.02	1	0	0
Q9DBZ1	Inhibitor of nuclear factor kappa-B kinase-interacting protein OS=Mus musculus GN=Ikbip PE=1	1.00	1.00	1.00	0.07	0	0	0
Q9DCF9	Translocon-associated protein subunit gamma OS=Mus musculus GN=Ssr3 PE=1 SV=1	1.00	1.00	1.00	0.02	0	0	0
Q9DCN2	NADH-cytochrome b5 reductase 3 OS=Mus musculus GN=Cyb5r3 PE=1 SV=3	1.00	0.04	0.00	0.00	0	1	3
Q9DCX2	ATP synthase subunit d, mitochondrial OS=Mus musculus GN=Atp5h PE=1 SV=3	1.00	1.00	0.02	1.00	0	0	1
Q9QYB5	Gamma-adducin OS=Mus musculus GN=Add3 PE=1 SV=2	1.00	1.00	1.00	0.00	0	0	0
Q9QYC0	Alpha-adducin OS=Mus musculus GN=Add1 PE=1 SV=2	1.00	1.00	0.00	0.00	0	0	2
Q9R0P6	Signal peptidase complex catalytic subunit SEC11A OS=Mus musculus GN=Sec11a PE=2 SV=1	1.00	0.03	1.00	1.00	0	1	0
Q9R0Q3	Transmembrane emp24 domain-containing protein 2 OS=Mus musculus GN=Tmed2 PE=1 SV=1	0.02	0.00	0.02	0.03	1	2	1
Q9WV80	Sorting nexin-1 OS=Mus musculus GN=Snx1 PE=1 SV=1	1.00	1.00	0.02	0.02	0	0	1
Q9Z0X1	Apoptosis-inducing factor 1, mitochondrial OS=Mus musculus GN=Aifm1 PE=1 SV=1	1.00	0.00	1.00	0.06	0	2	0

p14	Spectral Counts			p14	Normalized Spectral Counts				gene
	e15.5	e17.5	p2		e15.5	e17.5	p2	p14	
35	0	12	22	74	0.0000	0.0017	0.0062	0.0208	Spna2
1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Sec22b
2	0	0	0	7	0.0000	0.0000	0.0000	0.0020	Ptgis
0	0	7	2	0	0.0000	0.0010	0.0006	0.0000	Phb2
1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Myadm
2	0	2	0	4	0.0000	0.0003	0.0000	0.0011	Ddost
0	0	3	0	0	0.0000	0.0004	0.0000	0.0000	Pgrmc1
2	0	0	0	2	0.0000	0.0000	0.0000	0.0006	Entpd2
1	0	0	0	2	0.0000	0.0000	0.0000	0.0006	Copb2
1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Cyp8b1
2	2	35	15	4	0.0015	0.0049	0.0043	0.0011	Aldoa
0	0	1	2	0	0.0000	0.0001	0.0006	0.0000	Itga5
0	0	1	0	0	0.0000	0.0001	0.0000	0.0000	H2-Q9
8	3	21	15	16	0.0023	0.0030	0.0043	0.0045	Lmnb1
6	0	0	0	12	0.0000	0.0000	0.0000	0.0034	Tgm2
3	33	68	24	9	0.0255	0.0096	0.0068	0.0025	Marcks
8	0	8	0	14	0.0000	0.0011	0.0000	0.0039	Canx
0	0	4	0	0	0.0000	0.0006	0.0000	0.0000	Fkbp2
1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Aldh3a2
1	0	1	2	2	0.0000	0.0001	0.0006	0.0006	Capza1
1	0	1	1	2	0.0000	0.0001	0.0003	0.0006	Capzb
0	0	2	0	0	0.0000	0.0003	0.0000	0.0000	Cox7a2
1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Cyb5a
1	2	1	0	1	0.0015	0.0001	0.0000	0.0003	Ucp3
0	3	5	0	0	0.0023	0.0007	0.0000	0.0000	Drd2
2	0	1	0	4	0.0000	0.0001	0.0000	0.0011	Dad1
0	0	0	2	0	0.0000	0.0000	0.0006	0.0000	Pafah1b1
4	8	7	2	8	0.0062	0.0010	0.0006	0.0023	Eif5a
0	0	0	1	0	0.0000	0.0000	0.0003	0.0000	Phb
2	0	0	4	5	0.0000	0.0000	0.0011	0.0014	Slc9a3r1
2	0	0	2	3	0.0000	0.0000	0.0006	0.0008	Anpep
0	0	2	0	0	0.0000	0.0003	0.0000	0.0000	Atp5j
8	0	0	0	18	0.0000	0.0000	0.0000	0.0051	Jup
1	0	0	4	2	0.0000	0.0000	0.0011	0.0006	Rac2
0	0	2	5	0	0.0000	0.0003	0.0014	0.0000	Atp5i
0	2	16	12	0	0.0015	0.0023	0.0034	0.0000	Igf2r
0	0	2	0	0	0.0000	0.0003	0.0000	0.0000	Dlk1
1	0	0	0	2	0.0000	0.0000	0.0000	0.0006	Esyt1
0	0	1	0	0	0.0000	0.0001	0.0000	0.0000	Trpm3
1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Arcn1
0	0	0	1	0	0.0000	0.0000	0.0003	0.0000	Reep5
1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Vdac3
2	0	13	5	4	0.0000	0.0018	0.0014	0.0011	Vdac1
0	0	0	3	0	0.0000	0.0000	0.0009	0.0000	Dmbt1
2	0	1	5	3	0.0000	0.0001	0.0014	0.0008	Tmpo
0	0	2	0	0	0.0000	0.0003	0.0000	0.0000	Pde3b
1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Gdi2
27	0	1	23	57	0.0000	0.0001	0.0065	0.0161	Sptbn1
1	0	14	6	2	0.0000	0.0020	0.0017	0.0006	Selenbp2
0	0	1	0	0	0.0000	0.0001	0.0000	0.0000	Ccpq1
0	0	2	0	0	0.0000	0.0003	0.0000	0.0000	Cd34
2	0	0	0	2	0.0000	0.0000	0.0000	0.0006	Cltc
0	3	0	0	0	0.0023	0.0000	0.0000	0.0000	Nav3
0	0	0	2	0	0.0000	0.0000	0.0006	0.0000	Mtdh
3	0	0	0	7	0.0000	0.0000	0.0000	0.0020	Ehd2
1	0	0	0	2	0.0000	0.0000	0.0000	0.0006	Asph
1	0	0	2	2	0.0000	0.0000	0.0006	0.0006	Aifm2
1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Copa
1	3	3	6	2	0.0023	0.0004	0.0017	0.0006	Sdha
1	0	3	6	6	0.0000	0.0004	0.0017	0.0017	Ssspta
3	0	1	4	6	0.0000	0.0001	0.0011	0.0017	Rpn1
0	0	1	0	0	0.0000	0.0001	0.0000	0.0000	Ggt7
2	0	4	2	5	0.0000	0.0006	0.0006	0.0014	Tmed9
1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Dctn2
11	0	34	20	20	0.0000	0.0048	0.0057	0.0056	Rrbp1
0	2	2	0	0	0.0015	0.0003	0.0000	0.0000	Cox6c
0	0	0	1	0	0.0000	0.0000	0.0003	0.0000	Atp5f1
2	2	5	11	4	0.0015	0.0007	0.0031	0.0011	Ssr1
0	0	0	1	0	0.0000	0.0000	0.0003	0.0000	Uqcr1
1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Cmtm6
0	0	0	2	0	0.0000	0.0000	0.0006	0.0000	Sec13
0	9	8	1	0	0.0069	0.0011	0.0003	0.0000	Atp5d
1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Sec23b

1	0	3	5	1	0.0000	0.0004	0.0014	0.0003	Chmp4b
0	1	2	1	0	0.0008	0.0003	0.0003	0.0000	Uqcrc2
0	0	0	1	0	0.0000	0.0000	0.0003	0.0000	Plin3
1	3	0	0	2	0.0023	0.0000	0.0000	0.0006	Rpn2
1	0	0	0	1	0.0000	0.0000	0.0000	0.0003	Ikbip
1	0	0	0	2	0.0000	0.0000	0.0000	0.0006	Ssr3
4	0	2	13	8	0.0000	0.0003	0.0037	0.0023	Cyb5r3
0	0	0	8	0	0.0000	0.0000	0.0023	0.0000	Atp5h
3	0	0	0	5	0.0000	0.0000	0.0000	0.0014	Add3
2	0	0	5	4	0.0000	0.0000	0.0014	0.0011	Add1
0	0	1	0	0	0.0000	0.0001	0.0000	0.0000	Sec11a
1	1	7	2	2	0.0008	0.0010	0.0006	0.0006	Tmed2
1	0	0	5	1	0.0000	0.0000	0.0014	0.0003	Snx1
1	0	2	0	1	0.0000	0.0003	0.0000	0.0003	Aifm1

protein	annotation	FDR				Unique Peptides				Spectral
		e15.5	e17.5	p2	p14	e15.5	e17.5	p2	p14	e15.5
E9Q557	Desmoplakin OS=Mus musculus GN=Dsp PE=3 SV=1	1.00	0.00	0.00	0.00	0	2	3	19	0
P09055	Integrin beta-1 OS=Mus musculus GN=Itgb1 PE=1 SV=1	1.00	0.03	0.08	0.04	0	1	1	1	0
P11688	Integrin alpha-5 OS=Mus musculus GN=Itga5 PE=1 SV=3	1.00	0.04	0.02	1.00	0	1	1	0	0
P13020	Gelsolin OS=Mus musculus GN=Gsn PE=1 SV=3	0.05	0.00	0.00	0.00	1	3	4	3	2
P16045	Galectin-1 OS=Mus musculus GN=Lgals1 PE=1 SV=3	0.02	0.00	0.00	0.00	1	6	4	7	7
P19157	Glutathione S-transferase P 1 OS=Mus musculus GN=Gstp1 PE=1 SV=2	1.00	0.02	1.00	0.00	0	1	0	3	0
P21981	Protein-glutamine gamma-glutamyltransferase 2 OS=Mus musculus GN=Tgm2 PE=1 SV=4	1.00	1.00	1.00	0.00	0	0	0	6	0
P26039	Talin-1 OS=Mus musculus GN=Tln1 PE=1 SV=2	1.00	0.02	0.02	0.00	0	1	1	5	0
P26040	Ezrin OS=Mus musculus GN=Ezr PE=1 SV=3	1.00	0.02	0.02	0.00	0	1	1	2	0
P26041	Moesin OS=Mus musculus GN=Msn PE=1 SV=3	0.03	0.00	0.00	0.00	1	5	3	4	3
P28843	Dipeptidyl peptidase 4 OS=Mus musculus GN=Dpp4 PE=1 SV=3	1.00	1.00	1.00	0.02	0	0	0	1	0
P52793	Ephrin-A1 OS=Mus musculus GN=Efna1 PE=2 SV=1	1.00	0.07	1.00	1.00	0	1	0	0	0
P56480	ATP synthase subunit beta, mitochondrial OS=Mus musculus GN=Atp5b PE=1 SV=2	0.00	0.00	0.00	0.00	4	6	3	9	21
P60766	Cell division control protein 42 homolog OS=Mus musculus GN=Cdc42 PE=1 SV=2	1.00	0.00	0.02	1.00	0	2	1	0	0
P62141	Serine/threonine-protein phosphatase PP1-beta catalytic subunit OS=Mus musculus GN=Ppp1cb PE=1 SV=3	1.00	0.02	1.00	1.00	0	1	0	0	0
P97300	Neuroplastin OS=Mus musculus GN=Nptn PE=1 SV=3	1.00	1.00	0.07	0.08	0	0	1	1	0
P97350	Plakophilin-1 OS=Mus musculus GN=Pkp1 PE=1 SV=1	1.00	1.00	1.00	0.02	0	0	0	1	0
Q02257	Junction plakoglobin OS=Mus musculus GN=Jup PE=1 SV=3	1.00	1.00	1.00	0.00	0	0	0	8	0
Q04857	Collagen alpha-1(VI) chain OS=Mus musculus GN=Col6a1 PE=2 SV=1	1.00	1.00	0.02	0.04	0	0	1	1	0
Q64314	Hematopoietic progenitor cell antigen CD34 OS=Mus musculus GN=Cd34 PE=1 SV=1	1.00	0.07	1.00	1.00	0	1	0	0	0
Q64727	Vinculin OS=Mus musculus GN=Vcl PE=1 SV=4	1.00	0.02	0.00	0.00	0	1	4	8	0
Q8VDD5	Myosin-9 OS=Mus musculus GN=Myh9 PE=1 SV=4	1.00	1.00	0.00	0.00	0	0	3	15	0
Q9QU10	Transforming protein RhoA OS=Mus musculus GN=Rhoa PE=1 SV=1	1.00	0.04	0.00	0.02	0	1	2	1	0
Q9Z1T2	Thrombospondin-4 OS=Mus musculus GN=Thbs4 PE=2 SV=1	1.00	0.05	1.00	1.00	0	1	0	0	0

I Counts			Normalized Spectral Counts				gene
e17.5	p2	p14	e15.5	e17.5	p2	p14	
3	5	48	0.0000	0.0004	0.0014	0.0135	Dsp
2	1	1	0.0000	0.0003	0.0003	0.0003	Itgb1
1	2	0	0.0000	0.0001	0.0006	0.0000	Itga5
20	19	5	0.0015	0.0028	0.0054	0.0014	Gsn
37	23	40	0.0054	0.0052	0.0065	0.0113	Lgals1
1	0	7	0.0000	0.0001	0.0000	0.0020	Gstp1
0	0	12	0.0000	0.0000	0.0000	0.0034	Tgm2
2	1	9	0.0000	0.0003	0.0003	0.0025	Tln1
12	4	2	0.0000	0.0017	0.0011	0.0006	Ezr
18	11	7	0.0023	0.0025	0.0031	0.0020	Msn
0	0	2	0.0000	0.0000	0.0000	0.0006	Dpp4
1	0	0	0.0000	0.0001	0.0000	0.0000	Efna1
22	9	17	0.0162	0.0031	0.0026	0.0048	Atp5b
3	3	0	0.0000	0.0004	0.0009	0.0000	Cdc42
3	0	0	0.0000	0.0004	0.0000	0.0000	Ppp1cb
0	1	1	0.0000	0.0000	0.0003	0.0003	Nptn
0	0	2	0.0000	0.0000	0.0000	0.0006	Pkp1
0	0	18	0.0000	0.0000	0.0000	0.0051	Jup
0	2	1	0.0000	0.0000	0.0006	0.0003	Col6a1
2	0	0	0.0000	0.0003	0.0000	0.0000	Cd34
4	7	14	0.0000	0.0006	0.0020	0.0039	Vcl
0	7	26	0.0000	0.0000	0.0020	0.0073	Myh9
1	7	2	0.0000	0.0001	0.0020	0.0006	Rhoa
2	0	0	0.0000	0.0003	0.0000	0.0000	Thbs4

protein	annotation	FDR				Unique Peptides	
		e15.5	e17.5	p2	p14	e15.5	e17.5
O08553	Dihydropyrimidinase-related protein 2 OS=Mus musculus GN=Dpysl2 PE=1 SV=2	1.00	0.02	0.02	0.00	0	1
O35639	Annexin A3 OS=Mus musculus GN=Anxa3 PE=1 SV=4	1.00	1.00	1.00	0.00	0	0
O35855	Branched-chain-amino-acid aminotransferase, mitochondrial OS=Mus musculus GN=Bcat2 PE=2 SV=2	1.00	0.03	1.00	1.00	0	1
P07356	Annexin A2 OS=Mus musculus GN=Anxa2 PE=1 SV=2	0.00	0.00	0.00	0.00	2	6
P10107	Annexin A1 OS=Mus musculus GN=Anxa1 PE=1 SV=2	1.00	1.00	0.00	0.00	0	0
P26443	Glutamate dehydrogenase 1, mitochondrial OS=Mus musculus GN=Glud1 PE=1 SV=1	1.00	1.00	0.02	1.00	0	0
P34884	Macrophage migration inhibitory factor OS=Mus musculus GN=Mif PE=1 SV=2	0.04	0.02	1.00	1.00	1	1
P49817	Caveolin-1 OS=Mus musculus GN=Cav1 PE=1 SV=1	1.00	1.00	0.05	0.00	0	0
P52927	High mobility group protein HMGI-C OS=Mus musculus GN=Hmga2 PE=1 SV=1	0.02	0.05	1.00	1.00	1	1
P57759	Endoplasmic reticulum resident protein 29 OS=Mus musculus GN=Erp29 PE=1 SV=2	1.00	0.05	1.00	1.00	0	1
P61168	D(2) dopamine receptor OS=Mus musculus GN=Drd2 PE=2 SV=2	0.04	0.03	1.00	1.00	1	1
P63101	14-3-3 protein zeta/delta OS=Mus musculus GN=Ywhaz PE=1 SV=1	0.00	0.00	0.00	0.00	5	11
P70441	Na(+)/H(+) exchange regulatory cofactor NHE-RF1 OS=Mus musculus GN=Slc9a3r1 PE=1 SV=3	1.00	1.00	0.02	0.00	0	0
P97352	Protein S100-A13 OS=Mus musculus GN=S100a13 PE=1 SV=1	1.00	1.00	0.02	0.02	0	0
P97449	Aminopeptidase N OS=Mus musculus GN=Anpep PE=1 SV=4	1.00	1.00	0.03	0.00	0	0
Q00519	Xanthine dehydrogenase/oxidase OS=Mus musculus GN=Xdh PE=1 SV=5	1.00	1.00	1.00	0.02	0	0
Q3TJ91	Lethal(2) giant larvae protein homolog 2 OS=Mus musculus GN=Llg2 PE=2 SV=2	1.00	0.04	1.00	1.00	0	1
Q61409	cGMP-inhibited 3',5'-cyclic phosphodiesterase B OS=Mus musculus GN=Pde3b PE=1 SV=2	1.00	0.02	1.00	1.00	0	1
Q61425	Hydroxyacyl-coenzyme A dehydrogenase, mitochondrial OS=Mus musculus GN=Hadh PE=1 SV=2	1.00	0.00	1.00	1.00	0	2
Q61879	Myosin-10 OS=Mus musculus GN=Myh10 PE=1 SV=2	1.00	1.00	1.00	0.02	0	0
Q64285	Bile salt-activated lipase OS=Mus musculus GN=Cel PE=1 SV=1	1.00	0.00	0.02	0.00	0	6
Q91V41	Ras-related protein Rab-14 OS=Mus musculus GN=Rab14 PE=1 SV=3	1.00	0.02	1.00	1.00	0	1
Q99P72	Reticulon-4 OS=Mus musculus GN=Rtn4 PE=1 SV=2	1.00	0.02	1.00	0.00	0	1
Q9D0K2	Succinyl-CoA:3-ketoacid-coenzyme A transferase 1, mitochondrial OS=Mus musculus GN=Oxct1 PE=1 SV=1	1.00	0.02	1.00	1.00	0	1
Q9D1D4	Transmembrane emp24 domain-containing protein 10 OS=Mus musculus GN=Tmed10 PE=2 SV=1	1.00	0.00	1.00	0.00	0	2
Q9EPC2	Fibroblast growth factor 23 OS=Mus musculus GN=Fgf23 PE=1 SV=1	1.00	0.03	1.00	1.00	0	1
Q9JIW9	Ras-related protein Ral-B OS=Mus musculus GN=Ralb PE=2 SV=1	1.00	0.05	1.00	1.00	0	1

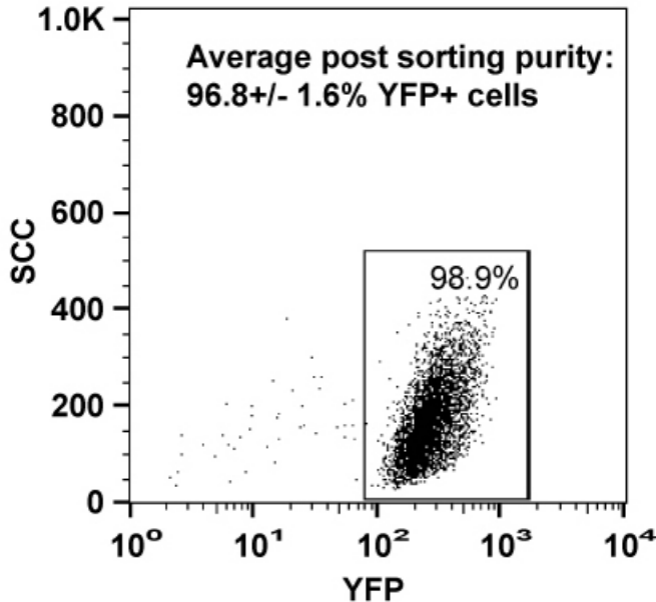
p2	Spectral Counts				Normalized Spectral Counts				gene	
	p14	e15.5	e17.5	p2	p14	e15.5	e17.5	p2		p14
1	7	0	14	6	12	0.0000	0.0020	0.0017	0.0033	Dpysl2
0	4	0	0	0	7	0.0000	0.0000	0.0000	0.0000	Anxa3
0	0	0	2	0	0	0.0000	0.0003	0.0000	0.0000	Bcat2
4	14	11	60	40	66	0.0085	0.0084	0.0114	0.0200	Anxa2
4	10	0	0	26	25	0.0000	0.0000	0.0074	0.0011	Anxa1
1	0	0	0	1	0	0.0000	0.0000	0.0003	0.0000	Glud1
0	0	2	2	0	0	0.0015	0.0003	0.0000	0.0000	Mif
1	2	0	0	2	9	0.0000	0.0000	0.0006	0.0000	Cav1
0	0	1	1	0	0	0.0008	0.0001	0.0000	0.0000	Hmga2
0	0	0	2	0	0	0.0000	0.0003	0.0000	0.0000	Erp29
0	0	3	5	0	0	0.0023	0.0007	0.0000	0.0000	Drd2
3	9	14	87	19	29	0.0108	0.0122	0.0054	0.0133	Ywhaz
1	2	0	0	4	5	0.0000	0.0000	0.0011	0.0000	Slc9a3r1
1	1	0	0	2	2	0.0000	0.0000	0.0006	0.0000	S100a13
1	2	0	0	2	3	0.0000	0.0000	0.0006	0.0000	Anpep
0	1	0	0	0	1	0.0000	0.0000	0.0000	0.0000	Xdh
0	0	0	2	0	0	0.0000	0.0003	0.0000	0.0000	Lig2
0	0	0	2	0	0	0.0000	0.0003	0.0000	0.0000	Pde3b
0	0	0	6	0	0	0.0000	0.0008	0.0000	0.0000	Hadh
0	1	0	0	0	2	0.0000	0.0000	0.0000	0.0000	Myh10
1	4	0	25	3	9	0.0000	0.0035	0.0009	0.0011	Cel
0	0	0	2	0	0	0.0000	0.0003	0.0000	0.0000	Rab14
0	3	0	1	0	4	0.0000	0.0001	0.0000	0.0000	Rtn4
0	0	0	2	0	0	0.0000	0.0003	0.0000	0.0011	Oxct1
0	3	0	4	0	4	0.0000	0.0006	0.0000	0.0000	Tmed10
0	0	0	2	0	0	0.0000	0.0003	0.0000	0.0000	Fgf23
0	0	0	1	0	0	0.0000	0.0001	0.0000	0.0000	Ralb

protein	annotation	FDR				Unique Peptides				Spectral
		e15.5	e17.5	p2	p14	e15.5	e17.5	p2	p14	e15.5
A2ASQ1	Agrin OS=Mus musculus GN=Agrn PE=2 SV=1	0.07	1.00	1.00	1.00	1	0	0	0	1
O55026	Ectonucleoside triphosphate diphosphohydrolase 2 OS=Mus musculus GN=Entpd2 PE=1 SV=2	1.00	1.00	1.00	0.00	0	0	0	2	0
P07356	Annexin A2 OS=Mus musculus GN=Anxa2 PE=1 SV=2	0.00	0.00	0.00	0.00	2	6	4	14	11
P07724	Serum albumin OS=Mus musculus GN=Alb PE=1 SV=3	1.00	0.02	1.00	0.00	0	1	0	4	0
P08228	Superoxide dismutase [Cu-Zn] OS=Mus musculus GN=Sod1 PE=1 SV=2	0.02	0.00	0.00	0.00	1	4	3	4	6
P09055	Integrin beta-1 OS=Mus musculus GN=Itgb1 PE=1 SV=1	1.00	0.03	0.08	0.04	0	1	1	1	0
P14211	Calreticulin OS=Mus musculus GN=Calr PE=1 SV=1	0.00	0.00	0.00	0.00	2	7	5	9	6
P16045	Galectin-1 OS=Mus musculus GN=Lgals1 PE=1 SV=3	0.02	0.00	0.00	0.00	1	6	4	7	7
P21981	Protein-glutamine gamma-glutamyltransferase 2 OS=Mus musculus GN=Tgm2 PE=1 SV=4	1.00	1.00	1.00	0.00	0	0	0	6	0
Q04857	Collagen alpha-1(VI) chain OS=Mus musculus GN=Col6a1 PE=2 SV=1	1.00	1.00	0.02	0.04	0	0	1	1	0
Q3V1T4	Prolyl 3-hydroxylase 1 OS=Mus musculus GN=Lepre1 PE=2 SV=2	1.00	1.00	1.00	0.02	0	0	0	1	0
Q60997	Deleted in malignant brain tumors 1 protein OS=Mus musculus GN=Dmbt1 PE=1 SV=2	1.00	1.00	0.00	1.00	0	0	2	0	0
Q9D1D6	Collagen triple helix repeat-containing protein 1 OS=Mus musculus GN=Cthrc1 PE=2 SV=1	1.00	0.05	1.00	1.00	0	1	0	0	0
Q9D8U4	Complement C1q tumor necrosis factor-related protein 2 OS=Mus musculus GN=C1qtnf2 PE=2 SV=1	1.00	0.02	0.03	1.00	0	1	1	0	0
Q9Z1T2	Thrombospondin-4 OS=Mus musculus GN=Thbs4 PE=2 SV=1	1.00	0.05	1.00	1.00	0	1	0	0	0

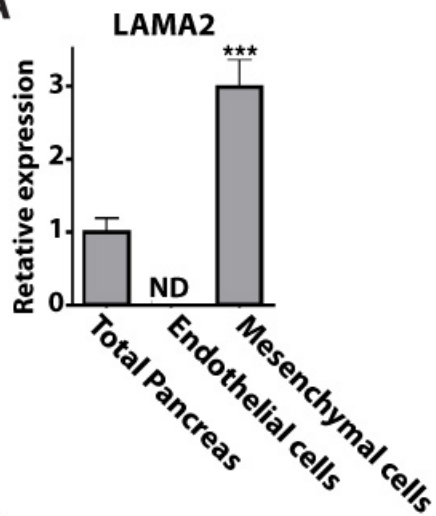
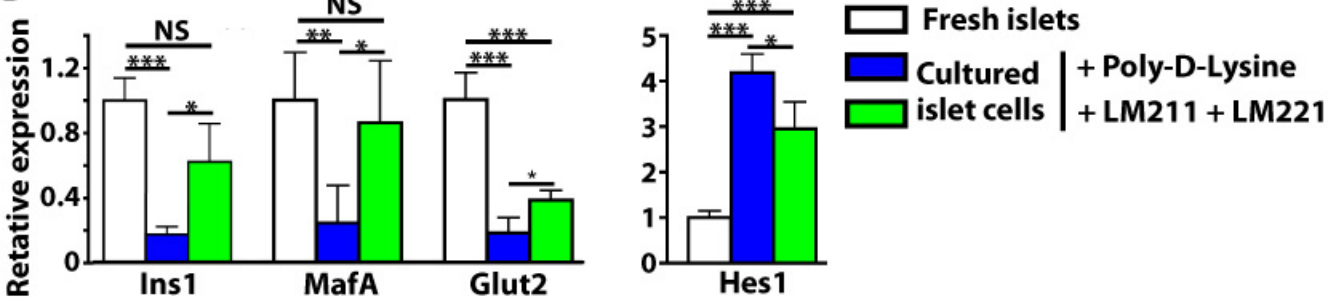
I Counts			Normalized Spectral Counts				gene
e17.5	p2	p14	e15.5	e17.5	p2	p14	
0	0	0	0.0008	0.0000	0.0000	0.0000	Agm
0	0	2	0.0000	0.0000	0.0000	0.0006	Entpd2
60	40	66	0.0085	0.0084	0.0114	0.0186	Anxa2
8	0	6	0.0000	0.0011	0.0000	0.0017	Alb
34	20	15	0.0046	0.0048	0.0057	0.0042	Sod1
2	1	1	0.0000	0.0003	0.0003	0.0003	Itgb1
53	21	22	0.0046	0.0075	0.0060	0.0062	Calr
37	23	40	0.0054	0.0052	0.0065	0.0113	Lgals1
0	0	12	0.0000	0.0000	0.0000	0.0034	Tgm2
0	2	1	0.0000	0.0000	0.0006	0.0003	Col6a1
0	0	1	0.0000	0.0000	0.0000	0.0003	Lepre1
0	3	0	0.0000	0.0000	0.0009	0.0000	Dmbt1
2	0	0	0.0000	0.0003	0.0000	0.0000	Cthrc1
4	2	0	0.0000	0.0006	0.0006	0.0000	C1qtnf2
2	0	0	0.0000	0.0003	0.0000	0.0000	Thbs4

protein	annotation	FDR				Unique Peptides		
		e15.5	e17.5	p2	p14	e15.5	e17.5	p2
P49817	Caveolin-1 OS=Mus musculus GN=Cav1 PE=1 SV=1	1.00	1.00	0.05	0.00	0	0	1
P60766	Cell division control protein 42 homolog OS=Mus musculus GN=Cdc42 PE=1 SV=2	1.00	0.00	0.02	1.00	0	2	1
P61168	D(2) dopamine receptor OS=Mus musculus GN=Drd2 PE=2 SV=2	0.04	0.03	1.00	1.00	1	1	0
P67871	Casein kinase II subunit beta OS=Mus musculus GN=Csnk2b PE=1 SV=1	1.00	1.00	0.06	1.00	0	0	1
P68040	Guanine nucleotide-binding protein subunit beta-2-like 1 OS=Mus musculus GN=Gnb2l1 PE=1 SV=3	1.00	1.00	0.02	0.05	0	0	1
P70441	Na(+)/H(+) exchange regulatory cofactor NHE-RF1 OS=Mus musculus GN=Slc9a3r1 PE=1 SV=3	1.00	1.00	0.02	0.00	0	0	1
Q3U1J4	DNA damage-binding protein 1 OS=Mus musculus GN=Ddb1 PE=1 SV=2	1.00	1.00	0.08	1.00	0	0	1
Q91X79	Elastase 1, pancreatic OS=Mus musculus GN=Cela1 PE=2 SV=1	1.00	0.00	0.02	0.00	0	4	1

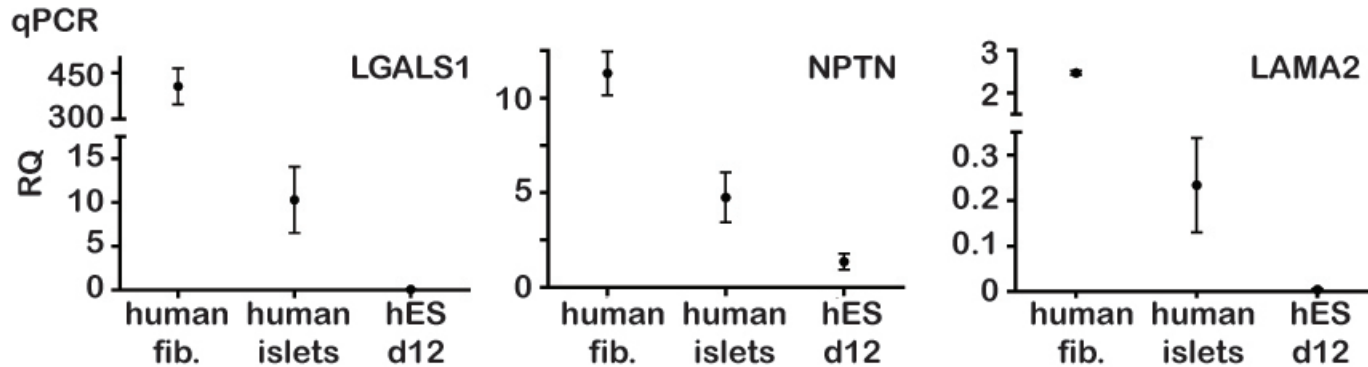
p14	Spectral Counts			p14	Normalized Spectral Counts				gene
	e15.5	e17.5	p2		e15.5	e17.5	p2	p14	
2	0	0	2	9	0	0	0.00057	0.00253	Cav1
0	0	3	3	0	0	0.00042	0.00085	0	Cdc42
0	3	5	0	0	0.00231	0.0007	0	0	Drd2
0	0	0	2	0	0	0	0.00057	0	Csnk2b
1	0	0	3	2	0	0	0.00085	0.00056	Gnb2l1
2	0	0	4	5	0	0	0.00114	0.00141	Slc9a3r1
0	0	0	1	0	0	0	0.00028	0	Ddb1
4	0	28	4	11	0	0.00394	0.00114	0.0031	Cela1



Supplementary Figure 1, Russ et al.

A**B**

Supplementary Figure 2, Russ et al.



Supplementary Figure 3, Russ et al.